

Discovery of reversible DNA methyltransferase and lysine methyltransferase G9a inhibitors with antitumoral in vivo efficacy

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J. Med. Chem., **Just Accepted Manuscript** • DOI: 10.1021/acs.jmedchem.7b01926 • Publication Date (Web): 28 Jun 2018

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3 **Discovery of reversible DNA methyltransferase and lysine**
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5 **methyltransferase G9a inhibitors with antitumoral in vivo efficacy**
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43

44 **ABSTRACT**

45
46 Using knowledge- and structure-based approaches, we designed and synthesized
47 reversible chemical probes that simultaneously inhibit the activity of two epigenetic
48 targets, histone 3 lysine 9 methyltransferase (G9a) and DNA methyltransferases
49 (DNMT), at nanomolar ranges. Enzymatic competition assays confirmed our design
50 strategy: substrate competitive inhibitors. Next, an initial exploration around our hit **11**
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3 was pursued to identify an adequate tool compound for *in vivo* testing. *In vitro* treatment
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5 of different hematological neoplasia cell lines led to the identification of molecules with
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7 clear anti-proliferative efficacies (GI_{50} values in the nanomolar range). Based on
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9 epigenetic functional cellular responses (levels of lysine 9 methylation and 5-
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11 methylcytosine), an acceptable therapeutic window (around 1 log unit) and a suitable
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13 pharmacokinetic profile, **12** was selected for *in vivo* proof-of-concept (ref 53). Herein,
14
15 **12** achieved a significant *in vivo* efficacy: 70% overall tumor growth inhibition of a
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17 human AML (Acute Myeloid Leukemia) xenograft in a mouse model.
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22 INTRODUCTION

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26 Epigenetic regulation of gene expression is controlled by DNA methylation,
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28 nucleosome positioning, histone variant exchange and post-translational histone
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30 modifications.^{1,2} Epigenetic alterations contribute to many human diseases, including
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32 cancer, inflammation, brain disorders, and metabolic and cardiovascular diseases. Thus,
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34 the development of epigenetic-based therapies has attracted considerable interest in the
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36 scientific and pharmaceutical communities, especially after the approval of the first
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38 therapies targeting DNA methyltransferases (DNMTs) and histone deacetylases
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40 (HDACs) for treatment of hematological malignancies. Thus, among possible
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42 epigenetic modifications, methylation has gained increased attention.
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44 Methyltransferases, the methyl writers of the histone and DNA code, catalyze the
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46 transfer of a methyl group from S-adenosyl-L-methionine (SAM or AdoMet) to its
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48 corresponding substrate—histones, nonhistone proteins or DNA—yielding S-adenosyl-
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50 L-homocysteine (SAH or AdoHcy). Methyltransferases acting on lysine and arginine
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3 residues of histones and other proteins are referred to as Protein Methyltransferases
4 (PMTs), while those that methylate DNA are known as DNMTs.

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7 G9a (also known as KMT1C (lysine methyltransferase 1C) or EHMT2 (euchromatic
8 histone methyltransferase 2)) is a histone-lysine N-methyltransferase that catalyzes the
9 transfer of one or two methyl groups to the ϵ -amino group of lysine 9 of histone H3
10 (H3K9me1 and H3K9me2), a hallmark associated with transcriptional gene silencing.
11
12 Other protein targets³ of G9a include the tumor suppressor p53, whose methylation leads
13 to its inactivation.⁴ G9a is upregulated in various cancers, and its overexpression has
14 been associated with poor prognosis and metastasis.⁵⁻⁷ There is mounting evidence that
15 the knockdown or pharmacological inhibition of G9a decreases cell growth,⁸⁻¹¹ delays
16 acute myeloid leukemia (AML) progression¹² and blocks tumor metastasis.^{6,7} Beyond
17 cancer, G9a is involved in the maintenance of HIV-1 latency,¹³ colitis,¹⁴ cognitive
18 disturbances (mental retardation, cocaine addiction, age-related cognitive decline),¹⁵
19 embryonic development¹⁶ and stem cell reprogramming, which has been used to
20 produce inducible pluripotent stem cells (iPSCs).¹⁷

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Compound **1** (BIX-01294, Chart 1) was the first selective inhibitor of G9a and GLP
(G9a-like protein, its closely related protein, also known as EHMT1), discovered in a
screening campaign.¹⁸ As shown in Chart 1, subsequent optimizations of **1** led to the
identification of a series of compounds possessing the same quinazoline core. Thus,
inhibitor **2** (UNC-0224), around two log units more potent than **1**, was the first G9a-
small molecule inhibitor co-crystallized with G9a, demonstrating that these quinazoline-
based inhibitors bind at the substrate peptide binding site and are not SAM-competitive
inhibitors.¹⁹ Chemical probe **3** (UNC-0638)²⁰ was the result of an extensive medicinal
chemistry effort^{19,21} to obtain more potency against G9a and cell-permeable quinazoline
derivatives. Finally, compound **4** (UNC-0642), has an optimized pharmacokinetic (PK)

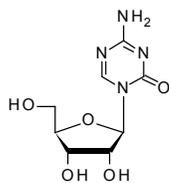
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3 profile for *in vivo* assays,²² and only very recently its efficacy in a Prader-Willi
4 syndrome mouse model has been proven.²³ Only a few studies have been published
5 aiming to identify new alternative chemotypes of the quinazoline ring: the chemical
6 probe **5** (A-366)²⁴ having a spiro[cyclobutane-1,3'-indole] ring and, more recently, the
7 2,4-diamino quinoline **6**, which was found to be a potent G9a inhibitor after exhaustive
8 heterocycle replacement of the quinazoline ring of BIX derivatives.²⁵ The *in vivo*
9 administration of compound **5** using osmotic mini-pump dosing in an AML flank
10 xenograft model resulted in moderate (45%) tumor growth inhibition.²⁶ With the
11 exception of **3** (IC₅₀ against DNMT1 in the micromolar range),^{20,27} all of these
12 compounds are inactive against DNMT1 at concentrations higher than 10 μM or even
13 50 μM. Other G9a inhibitors with a pharmacophore different from **1** have been
14 reviewed^{28–30} or were published very recently.^{31,32}

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29 The involvement of DNA methylation in cancer is well acknowledged.² DNMTs are
30 responsible for the methylation of the C5-cytosine of DNA. The methylation of
31 promoter CpG islands leads to gene inactivation, as described for tumor suppressors and
32 DNA repair genes in different tumors. DNMTs are grouped into two families: DNMT1
33 and DNMT3A/DNMT3B, each family being predominantly involved in methylation
34 maintenance (DNMT1) or de novo methylation (DNMT3A/DNMT3B).³³ These three
35 proteins are overexpressed in different cancer types or present mutations, as DNMT3A
36 in leukemia.³⁴ Additionally, there is recent evidence that DNA methylation may play an
37 important role in various diseases from different therapeutic areas: memory function,³⁵
38 Alzheimer's disease,³⁶ amyotrophic lateral sclerosis,³⁷ schizophrenia and bipolar
39 disorder,³⁸ addiction,³⁹ hypertension,⁴⁰ atherosclerosis,⁴¹ type 2 diabetes,⁴² systemic
40 lupus erythematosus⁴³ and fibrosis.⁴⁴ As mentioned above, two nucleoside analogs, **7**
41 (Azacytidine) and **8** (Decitabine) (Chart 2), have been approved by the U.S. FDA and
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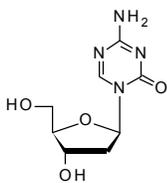
EMA for treatment of hematological malignancies. These drugs incorporate into DNA and inhibit DNMT1 through an irreversible covalent complex.⁴⁵ These two drugs show poor bioavailability, chemical instability and metabolic stability,⁴⁵ leading to the search for optimized nucleoside analogs, such as **9** (SGI-110), currently being explored in clinical trials.⁴⁶ However, due to the toxic side effects associated with its mechanism of action, there is increasing interest in finding potent non-incorporating reversible DNMT inhibitors. Among others (see refs^{47,48} for comprehensive reviews), the quinoline-based **10** (SGI-1027) has demonstrated inhibitory activity against DNMT1, DNMT3a and DNMT3b in biochemical assays and its potential to induce hypomethylation of CpG islands of tumor suppressor genes.⁴⁹ Interestingly, DNMT1 directly binds G9a and both methyltransferases cooperate during cell division,⁵⁰ promoting the transcriptional silencing of target genes,⁵¹ including the reactivation of tumor suppressor genes.⁵² Recently, we⁵³ and others⁵⁴ have shown that simultaneous blocking of G9a and DNMT1 exerts a synergistic effect on reducing the growth of cancer cells, either when combining G9a knockdown with pharmacologic inhibition of DNMT⁵⁴ or using specific siRNAs against both targets⁵³ or by inhibiting them with a combination of **5** and **8**.⁵³ Based on these results, we postulated that small-molecule reversible dual inhibitors of both methyltransferases might represent a novel approach in cancer therapeutics. Our efforts toward the identification of such dual inhibitors commenced with compound **11** (Chart 3), a quinoline-based matched pair of **3**. Subsequent optimization led to the discovery of the lead compound **12** (CM-272), with a suitable PK profile for *in vivo* administration and the ability to prolong the survival of AML, acute lymphoblastic leukemia (ALL) and diffuse large B-cell lymphoma (DLBCL) mice.⁵³ Here, we report the discovery and an initial exploration leading to the identification of our candidate for *in vivo* proof-of-concept compound **12**.

assay format.²⁷ The IC₅₀ values for compounds **2**¹⁹, **4**²², **5**²⁴, and **6**²⁵ have been published previously.

**7**

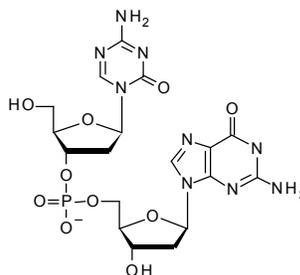
5-Azacididine

IC₅₀ G9a > 10000 nM
DNMT1 = 300 nM

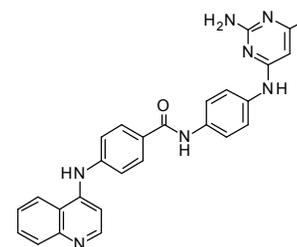
**8**

Decitabine

IC₅₀ G9a > 10000 nM
DNMT1 = 30 nM

**9**

SGI-110

**10**

SGI-1027

IC₅₀ G9a > 10000 nM
IC₅₀ DNMT1 = 3550 nM

Chart 2. Known DNMT1 inhibitors: nucleoside (compounds **7**, **8**, **9**) and non-nucleoside (compound **10**). G9a IC₅₀ values for **7**, **8** and **10** and DNMT1 IC₅₀ values for **10** as determined internally (see footnote in Table 1). Alternatively, DNMT1 IC₅₀ values of 12500 nM (Poly(dI-dC)) and 6000 nM (hemimethylated DNA) have been reported for compound **10** using a radioactive methyl transfer assay with two different substrates.⁴⁹ For G9a, reported value for **10** is 59 ± 4 μM.⁵⁷ DNMT1 values for **7** and **8** have been reported previously and correspond to the lower concentration at which DNA hypomethylation is observed.⁴⁵

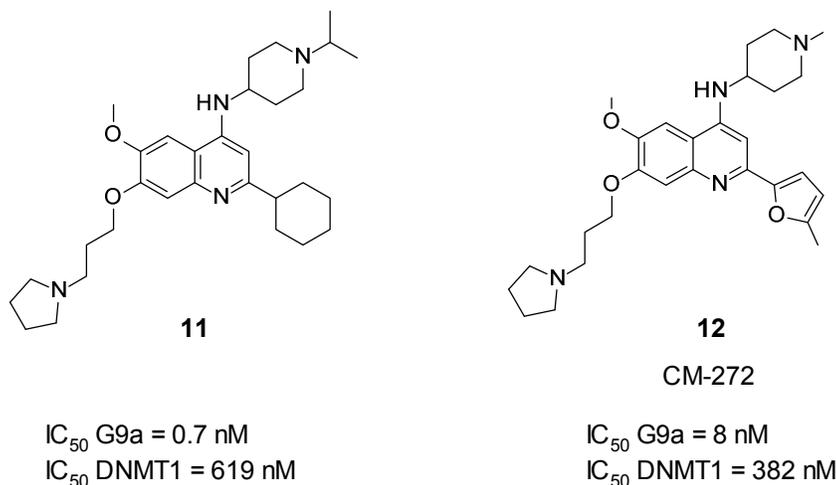


Chart 3. **11**, quinoline matched pair of **3**, and lead compound **12** (CM-272). G9a and DNMT1 IC_{50} values determined internally (see footnote in Table 1). Biochemical inhibitory data of **12**, and its close analogue (CM-579), was confirmed by orthogonal biochemical assays as well as by direct binding measurements to DNMT1 using microscale thermophoresis (MST) and by cellular functional response (using pyrosequencing).⁵³

RESULTS

Rational Design

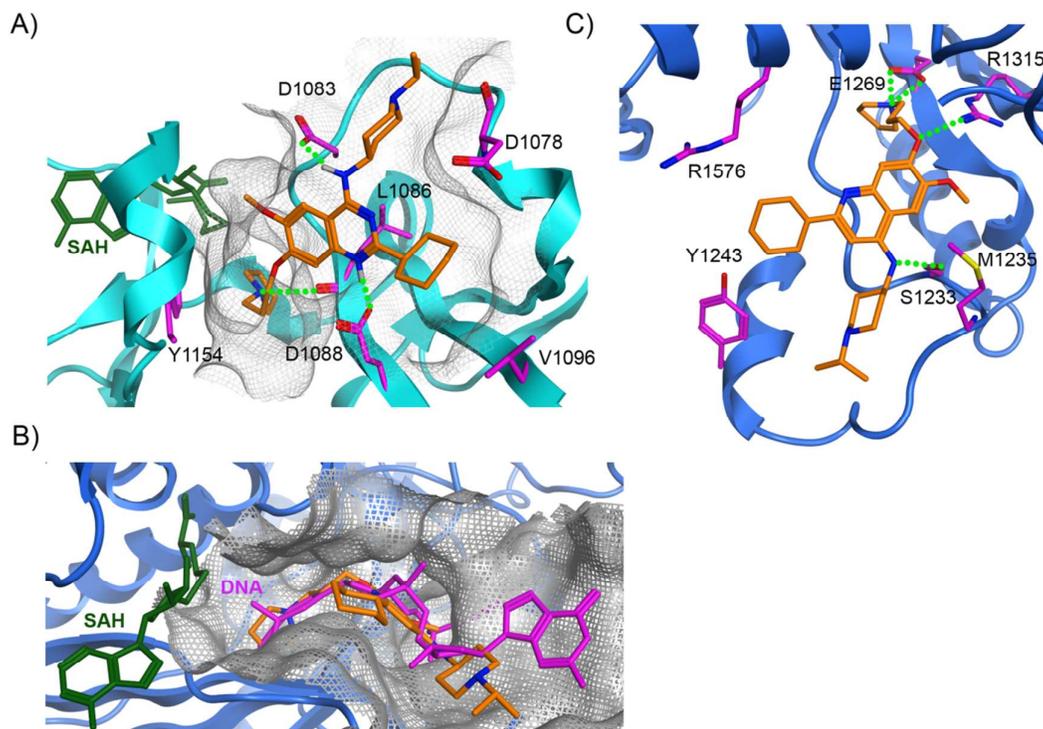
We pursued knowledge- and structure-based approaches to design novel G9a and DNMT methyltransferase activity inhibitors; targeting both enzymes. For G9a, the crystal structures of **2** (PDB entry 3K5K)¹⁹ and **3** (PDB entry 3RJW)²⁰ revealed that these quinazolinone-based inhibitors could bind to the histone-binding pocket of G9a.²⁰ A detailed analysis of the interactions of **3** with the substrate-binding pocket of G9a (Figure 1A) shows that i) the pyrrolidine side chain interacts with the lysine binding channel (hydrogen-bond contact with the carbonyl of Leu1086 and cation- π interaction with the side-chain of Tyr1154); ii) the -NH group at the 4-position of the quinazolinone

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3 establishes a hydrogen-bond with Asp1083; iii) the basic nitrogen of the piperidine ring,
4 which is directed toward the solvent region, is in close proximity with Asp1078 (4.5 Å);
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7 and iv) the N1 of the quinazoline ring, which is expected to be protonated at
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9 physiological pH,^{20,55} forms a salt bridge with Asp1088. As observed in Figure 1A, the
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11 ring nitrogen atom (N3) of the quinazoline scaffold does not participate in any direct
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13 key interaction with the protein, and no conserved water molecules appear proximal to
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15 this residue as mediators in a water-mediated hydrogen-bond contact. Thus, from the
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17 viewpoint of G9a activity, the replacement of the quinazoline ring by a quinoline would
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19 be well tolerated. Thus, we set out to synthesize and test the quinoline-matched pair of
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21 **3**, compound **11**, to validate our initial hypothesis for G9a activity. The observation that
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23 **3** displayed micromolar activity against DNMT1 (Chart 1) prompted us to use it as our
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25 initial starting point for the proof-of-concept of the dual G9a-DNMT inhibitor design.
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27 Moreover, during the course of our investigations, Rotili⁵⁶ disclosed a series of
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29 quinazoline-based compounds (based on chemical exploration around the G9a inhibitor
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31 **1**) that inhibit DNMT3A at low micromolar levels without any significant inhibition of
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33 DNMT1 and G9a.
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37 For the DNMT1 structure-based ligand design, different crystal structures of DNMT1 in
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39 complex with SAH, SAM, sinefungin (a SAM-competitive inhibitor) and/or DNA are
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41 publicly available. However, no crystal structures of any DNMT (including DNMT1,
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43 DNMT3A and DNMT3B) complexed with small molecule reversible inhibitors such as
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45 for example **10** are available, and only modeling studies have been published.⁵⁸⁻⁶¹ Thus,
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47 the binding mode as well as the binding cavity (SAM or DNA pockets), for most of
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49 these compounds, remain unknown. To guide the design of DNMT inhibitors, we
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51 carried out docking studies using the productive DNMT1-DNA-SAH complex structure
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53 (mouse DNMT1, mDNMT1, PDB entry 4DA4)⁶² containing a central hemimethylated
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3 CpG site and lacking the autoinhibitory linker (mDNMT1: 699-733) and zinc-finger-
4 like (CXXC motif, residues 646-692) domains at the N-terminal region of the catalytic
5 domain (mDNMT1, 1140-1602, hDNMT1 1139-1616). This structure was chosen
6 because the catalytic domain of the active DNMT1 undergoes a strong conformational
7 change compared with the autoinhibited structure,^{62,63} in which the DNA minor groove
8 is excluded from the catalytic site by the autoinhibitory linker. Thus, to model whether
9 compound **11** would bind to the SAM or the DNA pocket, a protein structure that can
10 accommodate both options was selected. Our docking studies of compound **11** showed
11 that the compound fits within the DNA binding pocket, superimposing well with the
12 hemimethylated CpG (Figure 1B) and the pyrrolidine ring occupying the cytidine-
13 binding pocket, establishing a hydrogen bond with the carboxylate group of the catalytic
14 Glu1269 (Glu1266 in human DNMT1, hDNMT1) (Figure 1C). Additionally, the
15 oxygen directly bonded at the 7-position establishes a hydrogen bond with the
16 guanidine of Arg1315 (Arg1312 in hDNMT1), a guanidine of the conserved motif TRR
17 flanking the entrance of the cytidine. The NH of the 4-position establishes a hydrogen
18 bond with the side chain of Ser1233 (Ser1230 in hDNMT1), and the isopropylpiperidyl
19 group overlays with the guanine of the DNA chain exiting the catalytic pocket and
20 establishing hydrophobic contacts with Met1235 (Met1232 in hDNMT1) (Figure 1C).
21 The methoxy group at the 6-position orients towards the 3' direction of the DNA chain.
22 Finally, the cyclohexyl ring at the 2-position lies in a cavity flanked by Thr1530
23 (Thr1528, hDNMT1), Gln1230 (Gln1227, hDNMT1), Tyr1243 (Tyr1240, hDNMT1)
24 and Arg1576 (Arg1574, hDNMT1), without making any explicit contact with any
25 residue. Comparing this active DNMT1 structure with the autoinhibited conformation of
26 DNMT1 (PDB entry 3PTA)⁶³ that has both the CXXC and autoinhibitory domains, it
27 can be seen that the C-terminal end of the CXXC linker (residues 692-700) lies
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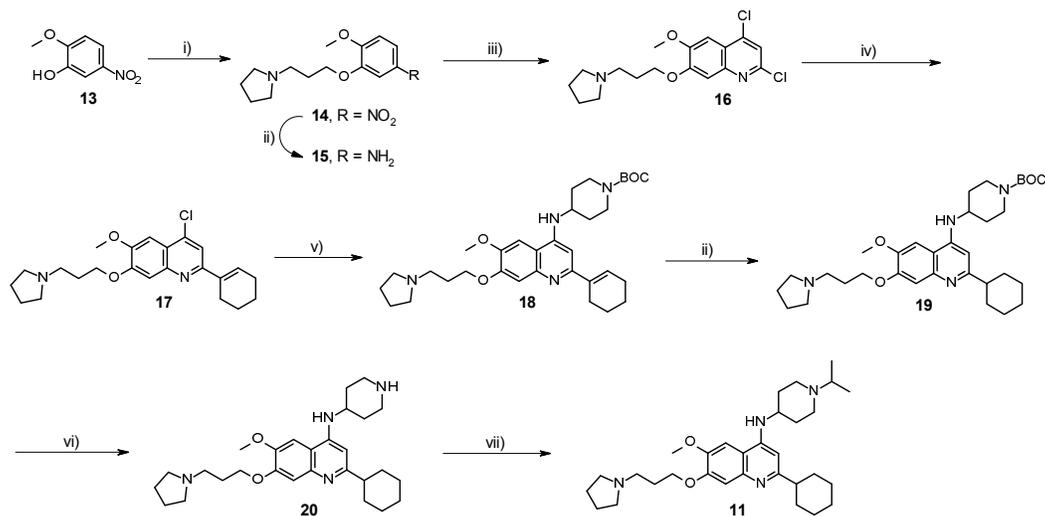
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3 proximal to the region occupied by the cyclohexyl group (Figure S1). Thus, without any
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5 further structural insight concerning the conformation adopted by the CXXC linker in
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7 the active form, no clear conclusion can be stated regarding the potential ligand
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9 interactions at this site. Similar conclusions were drawn when analyzing in detail a more
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11 recently released crystal structure of hDNMT1 with a higher resolution that is also auto-
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13 inhibited (PDB entry 4WXX).⁶⁴ Concerning the role of the central quinoline scaffold, it
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15 seems to act as a mere scaffold, without establishing any key contact with the protein.
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17 Thus, the replacement of the quinazoline of **3** by a quinoline **11** will predictably not
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19 have a strong influence on the inhibitory activity against DNMT1. In contrast to G9a,
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21 no apparent interaction between the nitrogen ring N1 and protein was observed, and no
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23 negatively charged residue lies in its neighborhood.
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53 **Figure 1.** (A) Complex of **3** with G9a (PDB accession code: 3RJW). (B) Docked
54 structure of compound **11** (orange) into DNMT1 (PDB accession code 4DA4) overlaid
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with DNA (pink). The cofactor SAH is shown in green. (C) Detailed analysis of the predicted hydrogen bonds between compound **11** and DNMT1.

Scheme 1.



Conditions: i) 3-pyrrolidin-1-yl-propan-1-ol, PPh_3 , DEAD, THF, 0 °C, followed by rt, 5 h; ii) Pd/C, MeOH, H_2 (1 atm), 15-25 °C, 3-10 h; iii) malonic acid, POCl_3 , rt, 4 h, followed by 90 °C overnight; iv) 2-(cyclohex-1-en-1-yl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane, K_2CO_3 , $\text{Pd}(\text{PPh}_3)_4$, 1,4-dioxane/ H_2O (5:1), MW, 120 °C, 1 h; v) *tert*-butyl 4-aminopiperidine-1-carboxylate, $\text{Pd}_2(\text{dba})_3$, BINAP, Cs_2CO_3 , 1,4-dioxane, 130 °C, 36 h; vi) HCl/EtOAc (1.0 M), 16 °C, 4 h; vii) acetone, AcOH, NaBH_3CN , THF, 50 °C, 15 h.

Compound **11** was synthesized as shown in Scheme 1 from commercially available 2-methoxy-5-nitrophenol (**13**). This compound was first transformed into amine (**15**) by the Mitsunobu reaction and hydrogenation. Subsequently, 2,4-dichloroquinoline (**16**) was prepared by reaction with malonic acid in POCl_3 , and intermediate **17** was isolated after Suzuki coupling. Next, the desired substitution at position 4 was installed by reaction with *tert*-butyl 4-aminopiperidine-1-carboxylate obtaining intermediate **18**, which was converted into compound **19** by hydrogenation. Finally, deprotection of the

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3 secondary amine under acidic conditions and reductive amination using acetone led us
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5 to the desired compound **11**.
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9 Compound **11** was evaluated for G9a and DNMT1 inhibitory activity using the time-
10 resolved fluorescence resonance energy transfer (TR-FRET) assay. Intriguingly,
11 compound **11** was more potent than **3** against the two targets, with IC₅₀ values of 0.7
12 nM against G9a (1.9 log units of increased potency) and 619 nM against DNMT1 (0.5
13 log units of increased potency) (Chart 3). Interestingly, in our hands, compound **3** was
14 also a better inhibitor of DNMT1 than compound **10** (Chart 2), a reversible DNMT
15 inhibitor commonly considered a reference compound for the DNMT inhibitor design.
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26 Concerning G9a activity, the higher potency of the quinoline than that of the
27 quinazoline scaffold is in line with the results recently published for the quinoline-
28 matched pair of compound **1** (BIX-01294), compound **6** (Chart 1).²⁵ In agreement with
29 the conclusions drawn by Srimongkolpithak et al, this improvement in G9a activity can
30 be attributed to the increased basicity of the N1 nitrogen that enhances the ionic
31 interaction with Asp1088, with the Pipeline Pilot⁶⁵ predicted pKa values of 6.54
32 (compound **11**) and 6.32 (compound **6**) for the quinoline series compared with the
33 predicted pKa values of 4.69 (compound **3**) and 4.48 (compound **1**) for their respective
34 quinazolines. Compared with the experimentally determined pKa values of N1 for
35 compound **6** (pKa of 9.50) and compound **1** (pKa of 6.94),²⁵ the pKa predictor
36 underestimates the pKa values, although the general trend is correctly predicted. Based
37 on the proposed binding mode of compound **11** into DNMT1, no explanation could be
38 stated for the improvement in the DNMT1 activity of compound **11** compared with that
39 of compound **3**.
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3 We next determined the mechanism of action of compound **11** by carrying out
4 enzymatic competition studies varying the concentrations of SAM and the two
5 substrates: histone monomethyl-H3K9 peptide for G9a and poly-deoxy inosine poly-
6 deoxy cytosine (pdI-pdC) DNA for DNMT1 (Figure 2). These competition experiments
7 confirmed that compound **11** is a substrate-competitive inhibitor of both targets, G9a
8 and DNMT1. In fact, the inhibitory potency of **11** is not affected by increasing
9 concentrations of the SAM cofactor (G9a IC_{50} ranging from 2 nM to 0.5 nM in Figure
10 2A and DNMT1 IC_{50} ranging from 1.4 μ M to 2.4 μ M in Figure 2C). However,
11 increasing the concentrations of the substrates, histone peptide (G9a, with IC_{50} values
12 spanning 0.7 nM to 29 nM, Figure 2B) and pdI-pdC (DNMT1, IC_{50} values from 1.4 μ M
13 to > 10 μ M, Figure 2D), have a clear impact on the binding of **11** to these two targets
14 (p IC_{50} differences are greater than 1 log unit). Non-competition with the SAM cofactor
15 is relevant to minimize off-target promiscuity. When compound **11** was assayed at a
16 concentration of 10 μ M against a panel of 14 lysine and arginine histone
17 methyltransferases, only GLP, a closely related protein to G9a, was considerably
18 inhibited (IC_{50} of 180 nM, with some selectivity towards G9a); the rest of histone
19 methyltransferases were marginally inhibited, lower than 25% inhibition
20 (Supplementary Table S1). For other DNMTs, only DNMT3A was significantly
21 inhibited by **11** at 10 μ M (80%), while had no effect on DNMT3B (Supplementary
22 Table S1).

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48 However, compound **11** had no anti-proliferative response against ALL CEMO-1 (GI_{50}
49 > 10 μ M), AML MV4-11 (GI_{50} > 10 μ M) and DLBCL OCI-Ly3 and OCI-Ly10 (0%
50 inhibition at 1 μ M) cell lines. Interestingly, it showed no cytotoxicity in the healthy
51 hepatic cell line THLE-2 (LC_{50} = 83.8 μ M).

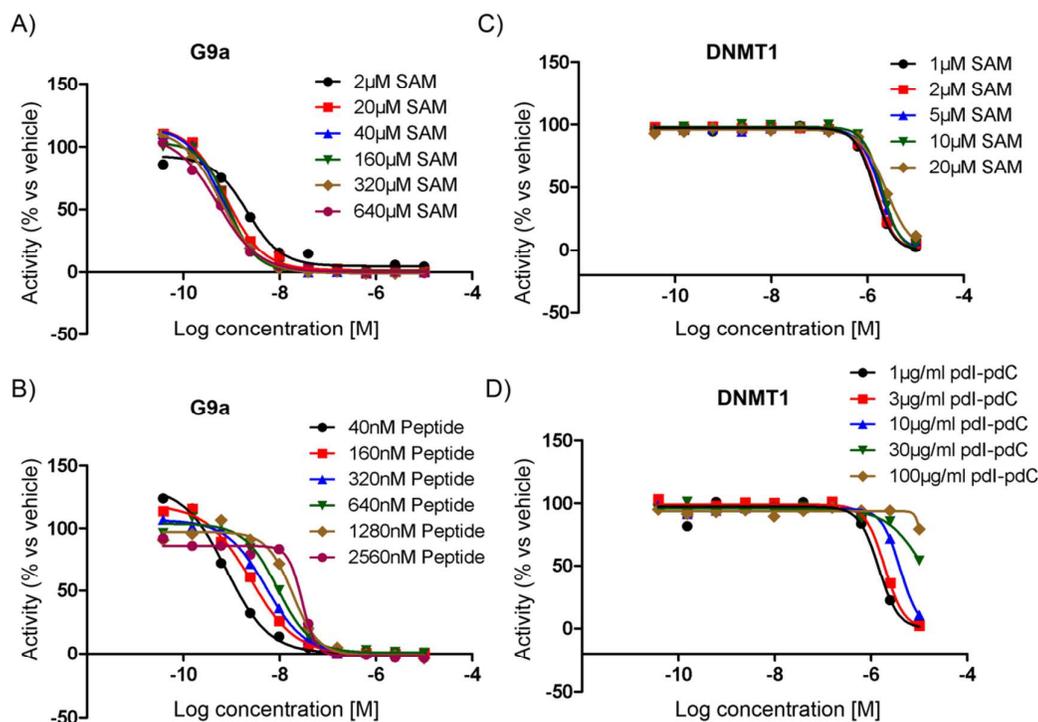


Figure 2. Competition experiments performed with compound **11** on G9a (A, B) and DNMT1 (C, D) by varying the SAM concentrations (A, C), histone monomethyl-H3K9 peptide (B) and pdI-pdC (C). Of note, the difference of 0.35 log units between the determined DNMT1 IC₅₀ value for this competition assay (1.4 μM) and the routinely screening (619 nM, Chart 3), was exceptionally accepted for the purpose of the competition assay (accepted difference in routinely screening is 0.3 log units, see footnote in Table 1; requirement is met by **11**).

In light of the high *in vitro* potency of **11** against G9a and DNMT1, we set out to optimize the quinoline series with the first initial goal of enhancing DNMT1 activity with potent anti-proliferative response against different hematological neoplasms and with an adequate ADME profile to be administered *in vivo* (Chart 4). Given, that GLP is

a closely related homologue of G9a, compounds were routinely screened against G9a, and only promising compounds were profiled against GLP.

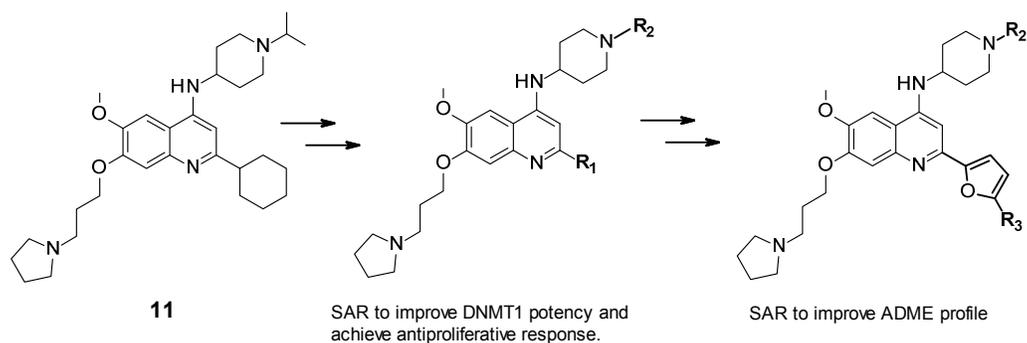


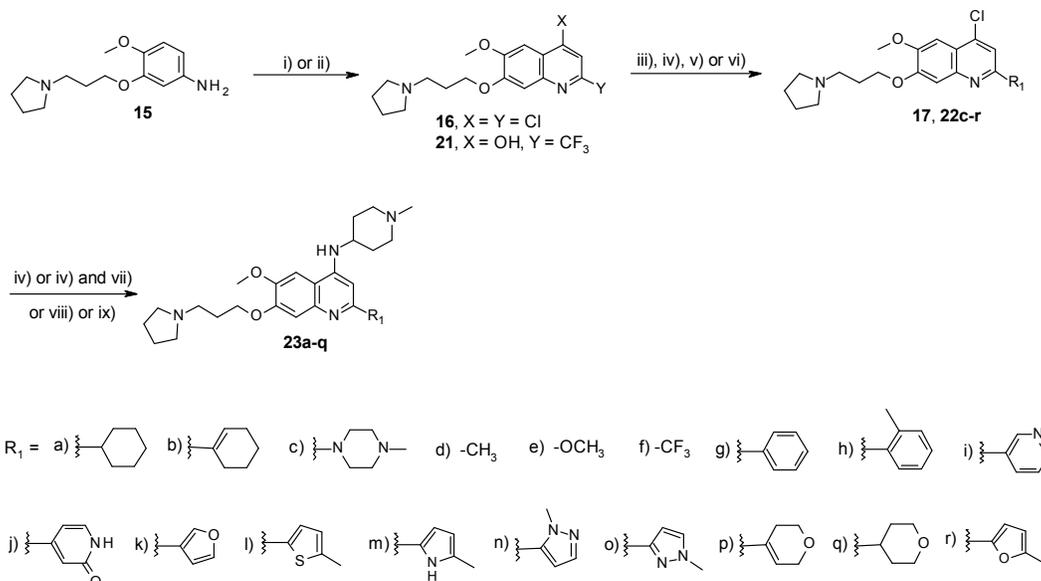
Chart 4. SAR exploration strategy to optimize our initial hit compound **11**.

SAR and antiproliferative response: Exploration of the 2-position (R_1).

Initially, we decided to focus our SAR exploration at the 2-position because i) modifications to the 2-substituent of the quinazoline scaffold are well tolerated from the viewpoint of G9a activity^{19,21,66}, and ii) we hypothesized that it might have a great impact on the permeability of the compounds to achieve a cellular response. For this exploration, the 1-methylpiperidin-4-ylamino substituent was fixed at the 4-position as the N-capping group because this fragment is directed toward the solvent-exposed region of both enzymes (predictably in the case of DNMT1), and it should not strongly influence the *in vitro* potency of compounds while reducing their molecular weight and easing synthetic accessibility. In fact, this fragment was found not to cause any potency loss against G9a compared with the 1-benzylpiperidin-4-ylamino group of compound **1** in the quinazoline series.¹⁹ Thus, compounds **23a-q** and **31** (Table 1) with diverse substituents at 2-position were designed and synthesized as shown in Schemes 2 and 3. The synthesis of compound **12** has been previously reported.⁵³ Compounds **23a-e** and

23g-q were prepared from intermediate **16**. First, the desired substitution at position 2 was installed by Suzuki coupling (**17**, **22d** and **22g-r**), Buchwald-Hartwig amination (**22c**) or reaction with NaOMe (**22e**). Next, the desired methylpiperidines were isolated by different Pd-mediated coupling reactions. In the case of compound **23f**, the trifluoromethyl group at position 2 was installed through reaction of intermediate **15** with 4,4,4-trifluoro-3-oxobutanoate, and then compound **22f** was prepared by heating in POCl₃. Finally, compound **23f** was synthesized by reaction with 1-methylpiperidin-4-amine under Buchwald-Hartwig amination conditions.

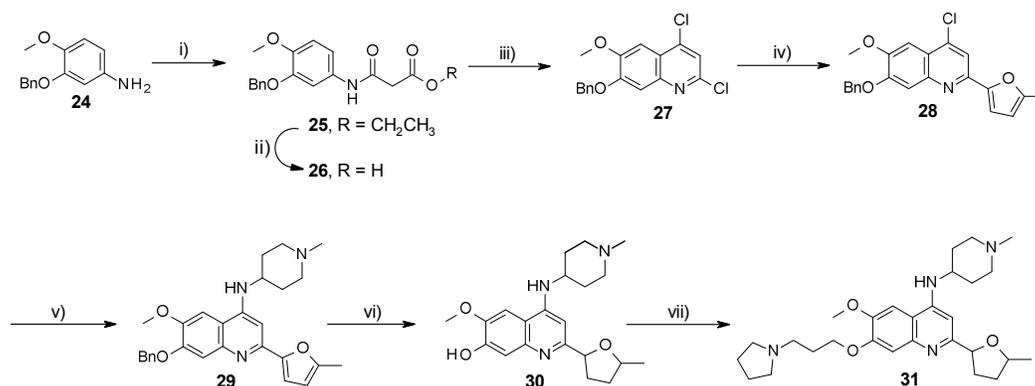
Scheme 2.



Conditions: i) malonic acid, POCl₃, rt, 4 h, followed by heating at 90 °C overnight; ii) 4,4,4-trifluoro-3-oxobutanoate, phenylpropanolamine, 120 °C, 12 h; iii) corresponding boronic acid or ester, K₂CO₃ or Na₂CO₃, Pd(PPh₃)₄, 1,4-dioxane or 1,4-dioxane/H₂O (5:1 or 15:1), 110-120 °C, MW or conventional heating, 1-12 h; iv) 1-methyl-piperazine or 1-methylpiperidin-4-amine, Pd₂(dba)₃, Cs₂CO₃, BINAP, 1,4-dioxane, 110-130 °C, MW or conventional heating, 1-12 h; v) NaOMe, rt, overnight; vi) POCl₃, 110 °C, 2 h; vii) Pd/C, EtOH, H₂ (1 atm), 25-35 °C, 4-15 h; viii) 1-methylpiperidin-4-amine, biphenyl-2-yl-

dicyclohexyl-phosphane, Pd₂(dba)₃, K₃PO₄, DME, 110 °C, MW, 3 h; ix) 1-methylpiperidin-4-amine, *t*-BuOK, xantphos, Pd₂(dba)₃, toluene, 130 °C, MW or conventional heating, 3-10 h.

Scheme 3.



Conditions: i) ethyl 3-chloro-3-oxo-propanoate, Et₃N, CH₂Cl₂, 0 °C, followed by rt, 12 h; ii) LiOH·H₂O, THF/MeOH/H₂O (3:3:2), rt, 16 h; iii) POCl₃, 90 °C, 2 h; iv) K₂CO₃, Pd(PPh₃)₄, 4,4,5,5-tetramethyl-2-(5-methyl-2-furyl)-1,3,2-dioxaborolane, 1,4-dioxane, 110 °C, 16 h; v) Cs₂CO₃, BINAP, Pd₂(dba)₃, 1-methylpiperidin-4-amine, 1,4-dioxane, 110 °C, 16 h; vi) Pd/C, H₂ (50 Psi), MeOH, 50 °C, 16 h; vii) Cs₂CO₃, 1-(3-chloropropyl)pyrrolidine, DMF, 110 °C, 16 h.

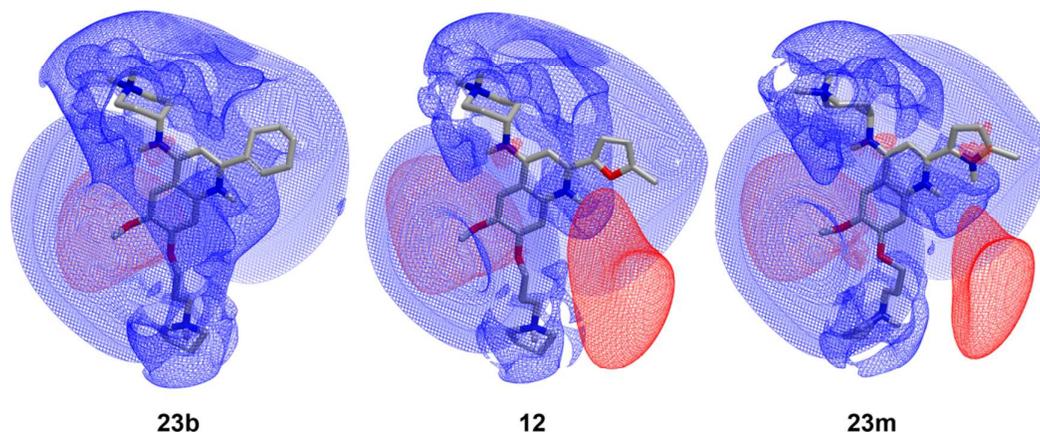
On the other hand, compound **31** (Scheme 3) was prepared from commercially available 3-benzyloxy-4-methoxy-aniline (**24**), which was converted into carboxylic acid **26** by reaction with ethyl 3-chloro-3-oxo-propanoate and ester hydrolysis. Next, 2,4-dichloroquinoline **27** was achieved after heating in POCl₃. Subsequent reaction with 4,4,5,5-tetramethyl-2-(5-methyl-2-furyl)-1,3,2-dioxaborolane led to intermediate **28**. As described above, the desired substitution at position 4 of the quinoline was then installed through Buchwald-Hartwig amination. Posterior hydrogenation afforded intermediate **30**, which was finally transformed into the desired compound **31** by reaction with 1-(3-chloropropyl)pyrrolidine in DMF.

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3 According to our initial hypothesis, the replacement of the N-capping group at the 4-
4 position of compound **11** had a minor influence on the *in vitro* potency against both
5 targets (compound **23a**, G9a and DNMT1 IC₅₀ values of 2 and 497 nM, respectively).
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9 From the viewpoint of G9a activity, in general, the cyclohexyl group of **23a** can be
10 replaced by diverse substituents (small groups, alkyl rings, aromatic rings,
11 heteroaromatic rings and heterocycles), most of them resulting in low nanomolar G9a
12 inhibitors (IC₅₀ < 50 nM). Analysis of the basicity of the N1 nitrogen can only partially
13 explain SAR results: decreased potency for electron-withdrawing groups such as –CF₃
14 (compound **23f**, 637 nM). However, it does not explain the reduced activity of other
15 small electron-donating groups such as the methoxy group of compound **23e** (136 nM).
16 Here, lipophilic interactions of 6-membered rings with Val1096 (Figure 1A) seem to
17 confer G9a potency, as solely an explicit potential hydrogen-bond interaction was
18 predicted for compound **23m** (between the NH of its methylpyrrole group and Asp1088,
19 Figure S2). Interestingly, this compound **23m** is the most potent against G9a in Table 1
20 (IC₅₀ = 0.5 nM). As for compound **11**, a substrate competitive mechanism was observed
21 for **23m** (Figure S3), indicating binding site conservation in both targets.
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39 Considering DNMT1 activity, a clear SAR trend was not observed for the diverse
40 substituents in Table 1. Small substituents such as methyl (**23d**), methoxy (**23e**) and –
41 CF₃ (**23f**) resulted in low micromolar (**23d**, **23e**) or inactive (IC₅₀ > 10 μM, **23f**)
42 compounds against DNMT1. Alkyl rings (**23a**, **23b**) tended to be more potent (IC₅₀
43 below or around 500 nM) than non-aromatic heterocycles such as N-methylpiperazine
44 (**23c**) and oxygen-containing rings (**31**, **23p**, **23q**) with DNMT1 inhibitory activity in
45 the low micromolar range (~ 1 μM) (Table 1). Regarding aromatic and heteroaromatic
46 rings, the introduction of 5-membered heteroaromatic rings such as 5-methyl-2-furyl
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3 (12), 5-methyl-2-thienyl (23i) and 5-methyl-1*H*-pyrrol-2-yl (23m) resulted in a slightly
4 increased DNMT1 inhibition compared with the cyclohexyl 23a (0.1-0.5 log units).
5 Analysis of the DNMT1 cavity around the 2-position of the quinoline core did not
6 reveal any clue to interpret the SAR, probably due to the missing CXXC domain.
7 However, comparative analysis of the electrostatic potential of these compounds
8 suggests a preference for negative potentials surrounding the 2-position of the
9 quinolines, as shown in Figure 3 for derivatives 12 (IC₅₀ = 382 nM) and 23m (IC₅₀ =
10 232 nM). However, as highlighted by compound 23b (IC₅₀ = 494 nM), this is not a strict
11 mandatory requirement.
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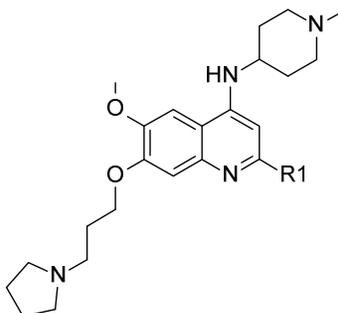
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Figure 3. Electrostatic maps of potent DNMT1 inhibitors 23b, 12 and 23m. The blue and red grids correspond to positive and negative potentials, respectively.

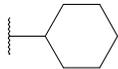
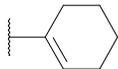
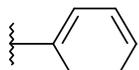
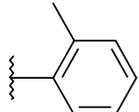
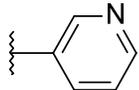
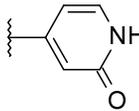
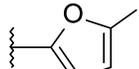
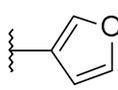
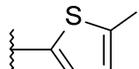
Concerning the anti-proliferative effect of the most biochemically potent compounds in Table 1 against ALL cell lines (CEMO-1 and primary culture LAL-CUN-2), DLBCL (OCI-Ly3 and OCI-Ly10) and AML (MV4-11) cell lines, only compound 12 showed GI₅₀ values lower than 1 μM against all five cell lines, indicating that 5-substituted-2-furyl groups at the 2-position are optimal for cellular activity. Other compounds with both, similar *in vitro* inhibitory profile and moderate PAMPA permeability (10 < Pe <

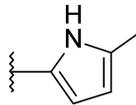
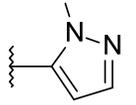
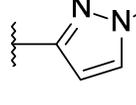
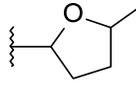
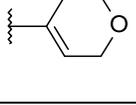
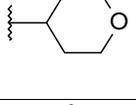
30 nm/s)⁶⁷ to that of **12**, had either no effect on proliferation (**23a**) or were poorer responders (**23b**, **23l**). Even compound **23m**, which displayed high activity in the *in vitro* assay against G9a and DNMT1, had minor effect on cellular viability than **12**. It remains to be seen if differences in active transport may account for this difference. For passive permeability, compounds with predicted LogD values at pH =7.4⁶⁵ ranging between 1 and 3 (**23a**, **23b**, **23g**, **12**, **23l**), exhibited better permeability (moderate) compared to more hydrophilic compounds (e.g. **23i**, **23o**, **23p**, **23q**), with Pe PAMPA values less than 10 nm/s). Cytotoxicity of selected compounds in the healthy hepatic cell line THLE-2 allowed for a therapeutic window (absolute difference between pLC₅₀ and pGI₅₀) of around 0.8 – 1 log units against their corresponding most responsive hematological neoplasia cell line (compounds **12**, **23b**, **23g**, **23l** and **23m**), or even higher (1.8 log units for compound **23k** in OCI-Ly10 cell line).

Table 1. Exploration of the 2-position



Cpd	R1	G9a	DNMT1	PAMP	CEMO-	LAL-	OCI-	OCI-	MV4-	THL
		IC ₅₀	IC ₅₀	A Pe	1	CUN-	Ly3	Ly10	11	E-2
		nM ^[e]	nM ^[e]	(nm/s)	GI ₅₀	2 GI ₅₀	GI ₅₀	GI ₅₀	GI ₅₀	LC ₅₀
				[d]	nM ^[e]	nM ^[e]	nM ^[e]	nM ^[e]	nM ^[e]	nM ^[f]

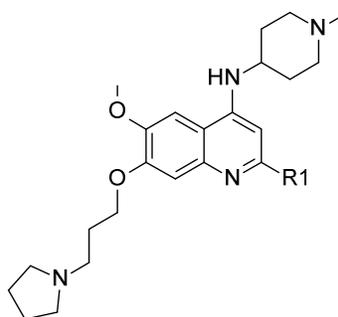
23a		2	497	19.8	20% ^[a]				>1000 0	34800
23b		6	494	20.9	1175	4290				7580
23c		42	945		14% ^[b]					
23d	Me	6	929		14% ^[b]					
23e	OMe	136	2290		1% ^[a]					
23f	CF ₃	637	>10000		24% ^[a]					
23g		15	1050	16.9	658	7% ^[a]	717	420	2595	5620
23h		52	582							
23i		40	1410	2.2	0% ^[a]	16% ^[a]	12% ^[a] 1	17% ^[a] 1		
23j		142	1250							
12 ^{3,5}		8	382	12.9	218	664	409	455	269	1780
23k		7	978	2.2	4560	54% ^[b]	>400 0	386		22300
23l		43	230	29.1	555		21% ^[a] 1	42% ^[a] 1	3095	4060

23m		0.5	232		1650	12% ^[a]	896	915		7680
23n		1080	>10000	0.9						
23o		44	969			10% ^[a]	16% ^[a]	16% ^[a]	0% ^[a]	
31 ^[g]		27	1370							
23p		12	957	3.7		8% ^[a]	0% ^[a]	6% ^[a]	16% ^[a]	
23q		17	4420	2.2		20% ^[a]	6% ^[a]	0% ^[a]	3% ^[a]	

^[a] Percentage of growth inhibition at 1 μ M of the compound; ^[b] percentage of growth inhibition at 10 μ M

of the compound. For data in Tables 1-3 and Charts 1-3: ^[c] all biochemical results are the average of at least two independent replicates performed at different days. If absolute pIC₅₀ difference was higher than 0.3 log units, additional replicates were performed until satisfying the experimental error (by discarding individual results with values outside 2 MADs of the mean value). ^[d] The PAMPA assay was performed in triplicate. ^[e] Proliferation assays are the average of three replicates at different days. ^[f] THLE-2 cytotoxicity results after 72 hours of incubation are the average of at least two independent experiments performed at different days. If absolute pLC₅₀ difference was higher than 1 log unit, additional replicates were performed until satisfying the experimental error (by discarding individual results with values outside 3 MADs of the mean value). ^[g] racemic mixture.

Subsequent SAR exploration of the 2-position was centered on studying the effects of the replacement of the methyl fragment of the 5-methyl-2-furyl group of compound **12**



Cp	R1	G9a IC ₅₀ nM ^[b]	DNMT 1 IC ₅₀ nM ^[b]	PAMP A Pe (nm/s)	CEMO -1 GI ₅₀ nM	LAL- CUN-2 GI ₅₀ nM	OCI- Ly3 GI ₅₀ nM	OCI- Ly10 GI ₅₀ nM
34a		2	234	23.9	56	577	95	64
34b		48	618	1.3	0% ^[a]	3% ^[a]	0% ^[a]	0% ^[a]
34c		148	1270					
34d		15	759	0.5	0% ^[a]	19% ^[a]	26% ^[a]	1% ^[a]
34e		264	1600					
34f		2720	1050					
36		3140	1070					

^[a] Percentage of growth inhibition at 1 μ M of the compound. For compound **34a**, GI₅₀ and LC₅₀ values in MV4-11 and THLE-2 cell lines are 679 nM and 2320 nM, respectively.

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3 Except for the 5-ethyl-2-furyl (compound **34a**), the introduction of different polar
4 substituents with either hydrogen bond donors (compounds **34b**, **34c**) or hydrogen bond
5 acceptors (compound **34e**) or both (compound **34d**) had a detrimental impact on both
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7 biochemical activities, emphasizing a stronger preference for hydrophobic groups.
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9 Bulky aromatic substituents (**34f**, **36**) were not well tolerated, especially for G9a,
10 probably caused by steric clashes with Val1096. Because no significant improvement
11 was observed, no further exploration was performed.
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18 Compound **34a**, with similar biochemical profile and slightly higher PAMPA
19 permeability than **12**, showed higher anti-tumour activity against all cell lines, except
20 for MV4-11. Especially, for CEMO-1, OCI-Ly3 and OCI-Ly10 lines ($GI_{50} < 100$ nM),
21 an interesting therapeutic window of 1.4 – 1.6 log units was achieved. In summary, as a
22 result of the exploration of the 2-position of the quinoline series, compounds **12** and **34a**
23 were selected for further profiling (see below).
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33 **SAR and anti-proliferative response: Exploration of the 4-position (R_2)**

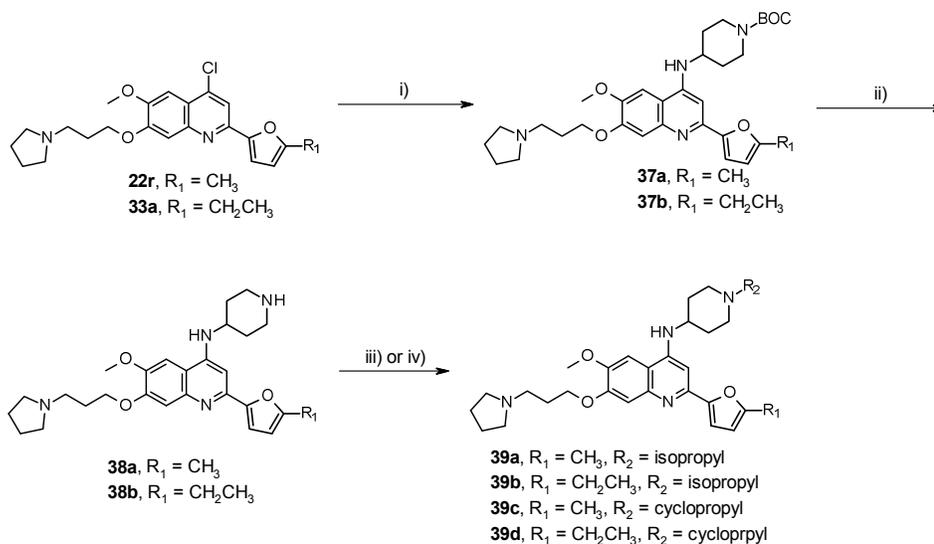
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38 Next, a small exploration of the N-capping group at the 4-position (cyclopropyl and
39 isopropyl groups) was carried out by holding constant the two optimal groups (5-
40 methyl-2-furyl and 5-ethyl-2-furyl) at the 2-position (Table 3). Our aim was two-fold:
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42 avoiding the potential N-demethylation *in vivo* of the 1-methylpiperidine ring and
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44 decreasing the basicity of the nitrogen of this piperidine to decrease potential hERG
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46 binding.
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53 Compounds **38a** and **39a-d** were prepared as illustrated in Scheme 5. Starting from
54 previously described 4-chloroquinolines **22r** and **33a**, BOC-protected piperidines **37a-b**
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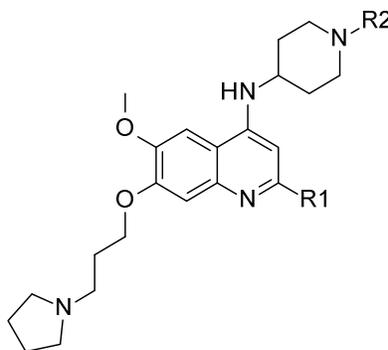
were isolated after Buchwald-Hartwig amination. Next, the BOC protecting group was removed under acidic conditions, and the desired isopropyl or cyclopropyl substituents were installed by reductive amination or cyclopropanation, respectively.

Scheme 5.



Conditions: i) *tert*-butyl 4-aminopiperidine-1-carboxylate, CS_2CO_3 , BINAP, $\text{Pd}_2(\text{dba})_3$, 1,4-dioxane, 80-130 °C, 5-16 h; ii) HCl/EtOAc (1.0 M), 22-25 °C, 2-3 h; iii) acetone, NaBH_3CN , AcOH, *i*-PrOH, 50 °C, 15 h; iv) (1-ethoxycyclopropoxy)-trimethyl-silane, NaBH_3CN , AcOH, MeOH, 60 °C, overnight.

Table 3. Exploration of the N-capping group at the 4-position



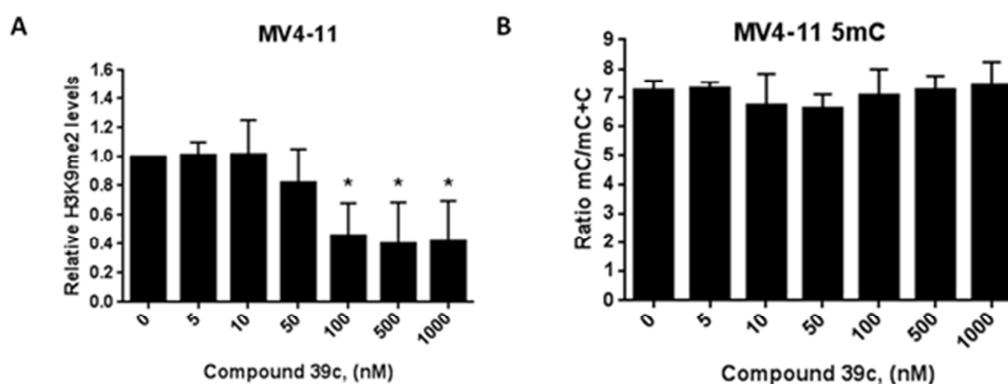
Cpd	R1	R2	G9a IC ₅₀ nM	DN MT1 IC ₅₀ nM	PAM PA Pe (nm/ s)	CE MO- 1 GI ₅₀ nM	LAL- CUN-2 GI ₅₀ nM	OCI- Ly3 GI ₅₀ nM	OCI- Ly10 GI ₅₀ nM	MV4 -11 GI ₅₀ nM	THL E-2 LC ₅₀ nM
38a		-H	6	137	2.0	1980	>5000	1270	>4000		8850
39a			3	231	25.8	247		898	792	649	1340
39b			3	284	38.3	441				971	1720
39c			22	413	40.4	453	1380	187	94	1590	455
39d			24	274	55.6	917		1560	236	3506	703

For DNMT1 activity, the replacement of the methyl group by an isopropyl or cyclopropyl group had a minor effect (IC₅₀ between 200 and 500 nM, as for **12** and **34a**). For G9a, the introduction of the cyclopropyl ring diminished G9a activity (up to 1 log unit in the case of compound **39d**), emphasizing the impact of the basicity of the alkylated nitrogen for G9a inhibitory activity, predictably decreased by the vinylic character of the cyclopropyl moiety.⁶⁹ Considering the trend in improved permeability (cyclopropyl > isopropyl > methyl), this drop in G9a potency might be one factor to explain the reduced cellular potency against CEMO-1, OCI-Ly3 and MV4-11 cell lines of compound **39d** (low micromolar range) compared to **34a** (low to middle nanomolar range). Notably, compound **39c** retained an acceptable growth inhibitory profile and was even more potent against DLBCL cell lines than compound **12**; then, **39c** was

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3 assayed at 10 μ M against a panel of 14 histone methyltransferases (Supplementary Table
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5 1), no target was significantly inhibited (>50%). Surprisingly, compound **39c** was
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7 selective for G9a over GLP (IC₅₀ > 20 μ M). For DNMTs, DNMT3A but not DNMT3B
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9 was inhibited by **39c** (81% and 0%, respectively). Thus, based on all these results, **39c**
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11 was further progressed into ADME and PK studies despite its THLE-2 cytotoxicity.
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13 Overall, these data showed that the inhibition of G9a and DNMT has differential
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15 phenotypic effects depending on the cell type. Despite a general trend in increased
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17 cytotoxicity in both tumoral and healthy cell lines can be observed (e.g. for compounds
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19 in Tables 2 and 3), this tendency was not confirmed for all molecules and, in some
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21 cases, adequate therapeutic windows are achieved (e.g. >1.5 log units for **23k** and **34a**).
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27 Finally, and as also reported for compound **12**,⁵³ the functional cellular potency of
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29 selected compounds was assessed by quantifying the global levels of H3K9me2
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31 (Western Blot) and 5-methylcytosine (5mC) (LC-MS/MS) in the MV4-11 cell line
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33 following 48 h of exposure. As an additional example of this chemical series, compound
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35 **39c** (Figure 4), with potent anti-proliferative effects (Table 3) and slightly worse
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37 biochemical profile than **12**, reduced significantly ($p \leq 0.05$) the H3K9me2 levels in a
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39 concentration-dependent manner from 50 nM (Figure 4A). In the case of the
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41 measurement of global DNA methylation by LC-MS/MS in the MV4-11 cell line, we
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43 detected that compound **39c** results in a subtle decrease, around 10%, of the overall
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45 DNA methylation in the genome at 50 nM (Figure 4B). Higher doses of **39c** did not
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47 result in reduced global DNA methylation levels; this seems to be a common fact to
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49 DNMT inhibitors Decitabine⁷⁰ and Azacitidine.⁷¹ In our opinion, these effects are
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51 because at high concentrations the majority of the cells are dead and we are only
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53 analyzing the epigenetic marks in those cells that are resistant to the treatment. This is a
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3 key point; we have to work at lower doses than their IC_{50} values, and long incubation
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5 periods, to avoid killing cells and thus to be able to monitor their impact on the
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7 epigenetic marks. To better verify the importance of the overall decrease in DNA
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9 methylation detected by LC-MS/MS, we conducted a study using MethylationEPIC
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11 BeadChip, which interrogate quantitatively over 850,000 methylation sites across the
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13 genome at single-nucleotide resolution. We performed this analysis in MV4-11 cells,
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15 controls as well as treated with 10nM and 50nM of **39c**. As identified by LC-MS/MS,
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17 both concentrations led to a subtle but very consistent change in the level of global
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19 DNA methylation (changes are greater than 5%); in fact, more than 15,000 CpGs are
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21 hypomethylated and there is a significant overlap between results obtained with both
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23 concentrations (Supplementary Information, Figure S4). The results obtained using LC-
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25 MS/MS and the methylation array show that the changes in the level of global and
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27 specific DNA methylation mediated by compound **39c** are small but in turn very
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29 consistent with all the techniques used and in all experiments carried out. In summary,
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31 these results demonstrate the functional dual effect of compound **39c** against the
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33 methyltransferase activity of G9a and DNMTs.
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51 **Figure 4.** H3K9me2 and 5mC hallmarks in the MV4-11 cell line after treatment with
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53 compound **39c** for 48 hours. A) Densitometry quantification of H3K9me2 levels. B)
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55 Global DNA methylation measurement by LC-MS/MS in MV4-11 cell line after
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3 treatment with different doses of compound **39c**. Asterisks show the P value of a one-
4
5 tailed Mann-Whitney U test (* $p \leq 0.05$). Error bars indicate s.d. from three replicates.
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8 9 **ADME and PK Profile**

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11 As shown in Table 4, the selected compounds **12**, **34a**, **39a** and **39c** exhibited low
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13 inhibition of the five major cytochrome P450 isoforms (1A2, 2C9, 2C19, 2D6, and
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15 3A4); in fact, the percentage inhibition values, at 10 μM , were less than 30% for all
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17 tested compounds. The metabolic stability of these three compounds at a concentration
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19 of 1 μM was evaluated in human and mouse liver microsomes after 20 min of
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21 incubation. A species difference in metabolism was observed. In human microsomes, 5-
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23 methyl-2-furyl substituted compounds **12** and **39c** displayed excellent stability (>95%
24
25 remaining), in contrast to the moderate stability of compound **34a** with the 5-ethyl-2-
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27 furyl side chain. In mice, only compound **12** had good stability (>70% remaining) and,
28
29 in contrast to our initial expectations, the introduction of the cyclopropyl ring of
30
31 compound **39c** did not improve its metabolic stability (<20% remaining), at least in
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33 mice.
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40 The low metabolic stability in mouse microsomes of compounds **34a** and **39c** translated
41
42 in a high clearance (**34a**) and a reduced half-life (**39c**, in comparison to **12**) when
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44 administered to BALB/c-RAG2^{-/-} $\gamma\text{C}^{-/-}$ mice for PK profiling (Table 5 and Supplementary
45
46 Tables S2 and S3); thus, precluding their use in *in vivo* efficacy assays. Only compound
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48 **12** dosed at 2.5 mg/kg intravenously (i.v.) revealed an acceptable profile with a good
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50 half-life (24.2 h) and an acceptable exposure ($\text{AUC}_{0-24\text{h}} = 1494 \text{ nM}$), optimal for once-
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52 daily dosing, to achieve *in vivo* efficacy *versus* cell lines with GI_{50} values around 200-
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54 450 nM.⁵³
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Table 4. ADME Profile of selected compounds

Cpd	1A2	2C9	2C19	2D6	3A4	HLM	HLM	MLM	MLM
	(%) ^a	(%) ^b	(t _{1/2}) ^c	(%) ^b	(t _{1/2}) ^c				
12⁵³	2.8	0.56	2.5	4.9	0.0	99.7	>2.4	70.9	0.6
34a	0.0	0.0	0.0	26.7	0.07	67.1	1.6	16.9	0.2
39c	28.6	8.0	4.0	27.6	4.6	97	>2.4	15	0.1
39a	4.3	6.1	0	11.4	0	64	1.7	36.9	0.3

^a % inhibition at 10 μM. ^b % compound remaining after a 20-min incubation in human or mouse liver microsomes (HLM and MLM respectively), ^c media time (h) in HML and MLM. All assays were performed in duplicate.

Table 5. PK Profile of selected compounds

Cpd	Route	Dose	AUC _{0-24h}	t _{1/2}	Cl	V _{ss}
		(mg/kg)	(nM * h)	(h)	(L/h/kg)	(L/kg)
12⁵³	i.v.	2.5	1494	24.2	0.91	29.3
34a	i.v.	1.9	505	13.2	5.90	88.5
39c	i.v.	5	8789	10.4	0.90	11.9

Species: BALB/c-RAG2^{-/-}γc^{-/-} mice; Vehicle: saline (NaCl 0.9%); n=4 and time points: 0.25, 1, 2, 4, 8 and 24 h (the time point after 1 h was not collected for compound **12**).

Compared with the PK profile of **4** (intraperitoneal administration, at 5mg/kg),²² the first G9a chemical probe suitable for *in vivo* testing, compound **12** (as reported in Table 5, i.v. at 2.5 mg/kg), reached a lower area under the curve (AUC_{0-24h}): 1494 nM versus 2314 nM, for **12** and **4**, respectively. Considering that intraperitoneal administration may lead to first pass clearance and the lower AUC_{0-24h} for **12**, one could assume (leaving aside the dose difference), that compound **4** exhibits a better PK profile than **12**. However, the once-daily administration of **12** at 2.5 mg/kg (i.v.) is optimal to reach

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3 an adequate sustained plasmatic concentration for *in vivo* treatment, in terms of efficacy
4 and safety (within the therapeutic window), and to achieve our initial aim: *in-vivo* proof-
5 of-concept (PoC). Additional medicinal chemistry efforts are required to optimize this
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7 molecule.
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10 11 12 13 ***In Vivo* Efficacy of CM-272 in human AML xenografts in a mouse model**

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15 Given the potent anti-proliferative response of compound **12** against the AML MV4-11
16 cell line, with a GI₅₀ value of 269 nM (Table 1),⁵³ we decided to test its *in vivo* efficacy
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18 in terms of tumor growth in a mouse model using human AML xenografts.
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24 This mouse model using human AML xenografts showed, for the first time, that tumor
25 growth is prevented using these reversible inhibitors, in tumors induced by
26 subcutaneously injecting 10×10⁶ MV4-11 cells in Rag2^{-/-}γc^{-/-} mice. These mice were
27 then treated with 2.5 mg/kg (i.v.) of compound **12** starting one day after leukemic cell
28 inoculation, administered daily during 21 consecutive days and sacrificed at day 23 after
29 cell inoculation. The mice were controlled for signs of morbidity (behavior and body
30 weight loss), and the tumor volume was monitored weekly (Figure 5). As shown in
31 Figure 5, treatment with **12** produced a significant (*p* value < 0.05) 70% overall tumor
32 growth inhibition (average tumor volumes at day 23 of 940 ± 435 mm³ and 3096 ± 1399
33 mm³ for treated and control groups, respectively). For comparison, when treating a
34 AML flank xenograft model using MV4-11 cells with compound **5**, administered via
35 osmotic mini-pump at 30 mg/kg/day for two weeks (to achieve a projected efficacious
36 exposure while mitigating the presumed C_{max}-driven toxicity), a modest 45% tumor
37 growth inhibition was observed.²⁶
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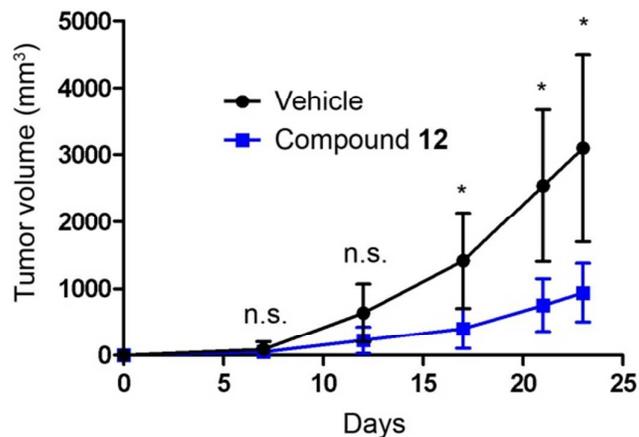


Figure 5. Compound **12** has *in vivo* activity in a MV4-11 tumor progression model. Average volume (mm³) \pm SD values of both groups, control and treated mice (n=6), during treatment. n.s. = non-statistically significant. * p value \leq 0.05.

DISCUSSION AND CONCLUSIONS

We presented a detailed account of the ligand- and structure-based design of a novel 4-aminoquinoline series of potent molecules that simultaneously inhibit G9a and DNMT1 methyltransferase activities in a reversible manner. Exploration around the 2- and 4-amino positions of the quinoline scaffold led us to evolve from the initial hit compound **11** and achieve a pharmacological tool compound, **12**, for *in vivo* proof-of-concept. Interestingly, the corresponding quinazoline-based pair of **12**, compound **40** (Chart 5 and synthesis in the Supporting Information), exhibited reduced biochemical potency, thereby validating the quinoline scaffold for the blocking of both targets, while quinazolines were extensively examined for G9a inhibition.

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3 selectivity profiling against the targets of interest (G9a and DNMTs: equally potent,
4 G9a selective and DNMT1 selective) has been also reported.⁷²
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9 **EXPERIMENTAL SECTION**

11 **Chemistry. General Procedure.**

13 Unless otherwise noted, all reagents and solvents were of the highest commercial
14 quality and used without further purification. All experiments dealing with moisture
15 sensitive compounds were conducted under N₂. Flash column chromatography was
16 performed on silica gel, particle size 60 Å, mesh = 230-400 (Merck) under standard
17 techniques. Automated flash column chromatography was performed using ready-to-
18 connect cartridges from Varian, on irregular silica gel, particle size 15-40 µm (normal
19 phase disposable flash columns) on a Biotage SPX flash purification system.
20 Microwave-assisted reactions were performed in a Biotage Smith Synthesis microwave
21 reactor. The NMR spectroscopic data were recorded on a Bruker AV400 or VARIAN
22 400MR spectrometer with standard pulse sequences, operating at 400 MHz. Chemical
23 shifts (δ) are reported in parts per million (ppm) downfield from tetramethylsilane
24 (TMS), which was used as internal standard. The abbreviations used to explain
25 multiplicities are s = singlet, d = doublet, t = triplet, m = multiplet. Coupling constants
26 (J) are in hertz. HPLC-analysis was performed using a Shimadzu LC-20AB or LC-
27 20AD with a Luna-C18(2), 5 µm, 2.0*50 mm column at 40 °C and UV detection at 215,
28 220 and 254 nm. Flow from the column was split to a MS spectrometer. The MS
29 detector (Agilent 1200, 6110MS or Agilent 1200, 6120MS Quadropole) was configured
30 with an electrospray source or API/APCI. N₂ was used as the nebulizer gas. The source
31 temperature was maintained at 50 °C. Data acquisition was accomplished with
32 ChemStation LC/MSD quad software. All tested compounds possessed a purity of at
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3 least 95% established by HPLC or LCMS unless otherwise noted. Reported yields were
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5 not optimized, the emphasis being on purity of product rather than quantity.
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10 **2-Cyclohexyl-*N*-(1-isopropyl-4-piperidyl)-6-methoxy-7-(3-pyrrolidin-1-
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12 ylpropoxy)quinolin-4-amine (11)**

13 To a mixture of compound **20** (90 mg, 0.19 mmol) and acetone (6 mg, 1.07 mmol) in
14 THF (30 mL) were added AcOH (65 mg, 1.08 mmol) and NaBH₃CN (67 mg, 1.07
15 mmol) in one portion at 16 °C under N₂ and the mixture was stirred at 50 °C for 15
16 hours. Then, the reaction mixture was cooled to 16 °C, filtered and concentrated in
17 vacuum. The residue was purified by preparative HPLC (method 1 described in
18 supporting information) to obtain pure compound **11** (38 mg, 39%) as a yellow syrup.
19
20 ¹H NMR (MeOD, 400 MHz): δ 7.81 (s, 1H), 7.31 (s, 1H), 6.75 (s, 1H), 4.37-4.30 (m,
21 2H), 4.27-4.21 (m, 1H), 4.03 (s, 3H), 3.87-3.80 (m, 2H), 3.64-3.55 (m, 3H), 3.52-3.45
22 (m, 2H), 3.37-3.28 (m, 2H), 3.17-3.07 (m, 2H), 2.91-2.82 (m, 1H), 2.41-2.34 (m, 4H),
23 2.22-1.92 (m, 10H), 1.86-1.68 (m, 4H), 1.56-1.46 (m, 2H), 1.42 (d, *J* = 8 Hz, 6H). ESI-
24 MS *m/z* 509.5 [M+H]⁺ calc. for C₃₁H₄₈N₄O₂.
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39 **1-[3-(2-Methoxy-5-nitro-phenoxy)propyl]pyrrolidine (14)**

40 To a solution of commercially available 2-methoxy-5-nitro-phenol (**13**) (49.0 g, 0.29
41 mol) in THF (500 mL) was added PPh₃ (152 g, 0.58 mol), 3-pyrrolidin-1-yl-propan-1-ol
42 (38 g, 0.29 mol) and DEAD (101 g 0.58 mol) at 0 °C and the solution was stirred at
43 room temperature for 5 hours. Then, the reaction mixture was concentrated and
44 extracted with EtOAc. The combined organic layers were washed with brine, dried over
45 anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
46 purified by column chromatography to obtain pure compound **14** (50 g, 62%) as yellow
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3 solid. ^1H NMR (CDCl_3 , 400 MHz): δ 7.88 (dd, $J = 8.8$ Hz, 1H), 7.76 (d, $J = 6.4$ Hz,
4 1H), 6.88 (d, $J = 8.8$ Hz, 1H), 4.16 (t, $J = 6.8$ Hz, 2H), 3.95 (s, 3H), 2.65 (t, $J = 7.2$ Hz,
5 2H), 2.60-2.45 (m, 4H), 2.10-2.04 (m, 2H), 1.85-1.75 (m, 4H). ESI-MS m/z 281
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7 $[\text{M}+\text{H}]^+$ calc. for $\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_4$.
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10 11 12 13 **4-Methoxy-3-(3-pyrrolidin-1-ylpropoxy)aniline (15)**

14
15 To a solution of compound **14** (14 g, 0.05 mol) in MeOH (200 mL) was added Pd/C (3
16 g) and the solution was stirred at 25 °C for 3 hours under H_2 atmosphere (1 atm). Then,
17 the mixture was filtrated and concentrated to give compound **15** (12 g, 96%) as yellow
18 oil. ^1H NMR (CDCl_3 , 400 MHz): δ 6.69 (d, $J = 8.4$ Hz, 1H), 6.33 (d, $J = 6.4$ Hz, 1H),
19 6.21 (dd, $J = 8.4$ Hz, 1H), 4.02 (t, $J = 6.8$ Hz, 2H), 3.76 (s, 3H), 2.65 (t, $J = 7.2$ Hz, 2H),
20 2.60-2.45 (m, 4H), 2.05-2.00 (m, 2H), 1.75-1.60 (m, 4H). ESI-MS m/z 251 $[\text{M}+\text{H}]^+$
21
22 calc. for $\text{C}_{14}\text{H}_{22}\text{N}_2\text{O}_2$.
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33 **2,4-Dichloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)quinoline (16)**

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35 To a solution of compound **15** (12.4 g, 0.049 mol) in POCl_3 (200 mL) was added
36 malonic acid (5.67 g, 0.055 mol) at room temperature. After stirring for 4 hours, the
37 solution was heated to 90 °C overnight. Then, the solution was concentrated and the
38 residue was poured into ice-water. The mixture was extracted with EtOAc and the
39 combined organic layers were washed with brine, dried over anhydrous Na_2SO_4 , filtered
40 and concentrated to give compound **16** (10 g, 57%) as pale yellow solid. ^1H NMR
41 (CDCl_3 , 400 MHz): δ 7.36 (s, 1H), 7.35 (s, 1H), 7.34 (s, 1H) 4.24 (t, $J = 6.8$ Hz, 2H),
42 4.02 (s, 3H), 2.66 (t, $J = 7.2$ Hz, 2H), 2.60-2.45 (m, 4H), 2.14 (t, $J = 6.8$ Hz, 2H), 1.87-
43 1.75 (m, 4H). ESI-MS m/z 355 $[\text{M}+\text{H}]^+$ calc. for $\text{C}_{17}\text{H}_{20}\text{Cl}_2\text{N}_2\text{O}_2$.
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4-Chloro-2-(cyclohexen-1-yl)-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)quinoline**(17)**

To a solution of compound **16** (708 mg, 2 mmol) in 1,4-dioxane/H₂O (5:1, 18 mL) was added K₂CO₃ (27 mg, 0.20 mmol), Pd(PPh₃)₄ (233 mg, 0.2 mmol) and 2-(cyclohex-1-en-1-yl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane (416 mg, 2 mmol) and the solution was heated to 120 °C under microwave irradiation for 1 hour. Then, the mixture was concentrated and extracted with EtOAc. The combined organic layers were washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by column chromatography to obtain pure compound **17** (0.4 g, 50%) as yellow solid. ESI-MS *m/z* 401 [M+H]⁺ calc. for C₂₃H₂₉ClN₂O₂. This intermediate was used in the next step without further characterization.

***Tert*-butyl 4-[[2-(cyclohexen-1-yl)-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-4-quinolyl]amino]piperidine-1-carboxylate (18)**

To a solution of compound **17** (150 mg, 0.38 mmol) and *tert*-butyl 4-aminopiperidine-1-carboxylate (375 mg, 1.87 mmol) in 1,4-dioxane (30 mL) were successively added Pd₂(dba)₃ (68 mg, 0.074 mmol), BINAP (93 mg, 0.15 mmol) and Cs₂CO₃ (305 mg, 0.94 mmol) and the resulting mixture was stirred at 130 °C for 36 hours under N₂. Then, the mixture was diluted with water and extracted with EtOAc. The combined organic phase was washed with brine, dried with Na₂SO₄, filtered, concentrated and purified by preparative TLC to give pure compound **18** (156 mg, 73%) as a yellow solid. ESI-MS *m/z* 565.4 [M+H]⁺ calc. for C₃₃H₄₈N₄O₄. This intermediate was used in the next step without further characterization.

***Tert*-butyl 4-[[2-cyclohexyl-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-4-quinolyl]amino]piperidine-1-carboxylate (19)**

A stirred suspension of compound **18** (156 mg, 0.28 mmol) in absolute MeOH (30 mL) containing Pd/C (250 mg) was placed under H₂ (15 Psi) at 15 °C for 10 hours. Then, the mixture was filtered through Celite and washed with MeOH (30 mL). The filtrate was concentrated to dryness to give compound **19** (158 mg, 99% crude) as a yellow solid. ESI-MS *m/z* 567.5 [M+H]⁺ calc. for C₃₃H₅₀N₄O₄. This intermediate was used in the next step without further characterization.

2-Cyclohexyl-6-methoxy-*N*-(4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (20)

A solution of compound **19** (95 mg, 0.17 mmol) in HCl/EtOAc (1.0 M, 15 mL) was stirred at 16 °C for 4 hours. Then, the reaction mixture was concentrated to dryness to give compound **20** (90 mg, 99% crude) as a yellow solid. ESI-MS *m/z* 467.4 [M+H]⁺ calc. for C₂₈H₄₂N₄O₂. This intermediate was used in the next step without further characterization.

6-Methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-(trifluoromethyl)quinolin-4-ol (21)

Phenylpropanolamine (50 mL) was heated to 80 °C with stirring in round-bottomed flask, then compound **15** (5 g, 0.02 mol) was added at 80-100 °C. After addition, ethyl 4,4,4-trifluoro-3-oxobutanoate (3.68 g, 0.02 mol) was added into the reaction mixture over 15-20 minutes. The reaction mixture was stirred vigorously at 120 °C for 12 hours. The reaction mixture was poured into ice-water and adjusted pH to 8 by Na₂CO₃, then concentrated and extracted with CH₂Cl₂/MeOH (3:1). The combined organic layer was concentrated to give the desired compound **21** (2.5 g, 33%). ESI-MS *m/z* 371 [M+H]⁺

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3 calc. for $C_{18}H_{21}F_3N_2O_3$. This intermediate was used in the next step without further
4
5 characterization.

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9 **4-Chloro-6-methoxy-2-(4-methylpiperazin-1-yl)-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22c)**

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12 To a solution of compound **16** (1.38 g, 3.9 mmol) in 1,4-dioxane (15 mL) was added
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14 Cs_2CO_3 (2.56 g, 7.86 mmol), BINAP (0.244 g, 0.39 mmol), $Pd_2(dba)_3$ (0.302 g, 0.33
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16 mmol) and 1-methyl-piperazine (0.6 g, 5.3 mmol) and the mixture was heated to 110 °C
17
18 for 12 hours. Then, the solution was concentrated and extracted with EtOAc. The
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20 combined organic layers were washed with brine, dried over anhydrous Na_2SO_4 , filtered
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22 and concentrated to give the crude product which was purified by column
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24 chromatography to obtain pure compound **22c** (0.3 g, 18%) as a yellow solid. ESI-MS
25
26 m/z 419 $[M+H]^+$ calc. for $C_{22}H_{31}ClN_4O_2$. This intermediate was used in the next step
27
28 without further characterization.
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35 **4-Chloro-6-methoxy-2-methyl-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22d)**

36
37 To a solution of compound **16** (2.78 g, 7.87 mmol) in 1,4-dioxane (30 mL) was added
38
39 $MeB(OH)_2$ (0.52 g, 8.66 mmol), K_2CO_3 (2.17 g, 15.7 mmol) and $Pd(PPh_3)_4$ (0.91 g, 0.78
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41 mmol) and the mixture was heated to 110 °C for 12 hours. Then, the solution was
42
43 concentrated and extracted with EtOAc. The combined organic layers were washed with
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45 brine, dried over anhydrous Na_2SO_4 , filtered and concentrated to give the crude product
46
47 which was purified by column chromatography to obtain pure compound **22d** (0.5 g,
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49 19%) as yellow solid. 1H NMR ($CDCl_3$, 400 MHz): δ 7.27 (s, 1H), 7.16 (s, 1H), 7.14 (s,
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51 19%) as yellow solid. 1H NMR ($CDCl_3$, 400 MHz): δ 7.27 (s, 1H), 7.16 (s, 1H), 7.14 (s,
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53 1H), 4.15 (t, $J = 6.8$ Hz, 2H), 3.92 (s, 3H), 2.60-2.51 (m, 5H), 2.50-2.41 (m, 4H), 2.10-
54
55 2.01 (m, 2H) 1.75-1.65 (m, 4H). ESI-MS m/z 335 $[M+H]^+$ calc. for $C_{18}H_{23}ClN_2O_2$.
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4-Chloro-2,6-dimethoxy-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22e)

Compound **16** (1.5 g, 4.24 mmol) was dissolved in NaOMe (25 mL, 25%) and stirred at room temperature overnight. Then, the reaction mixture was quenched with water and extracted with EtOAc. The combined organic phase was concentrated to give the crude product which was purified by preparative HPLC (method 1 described in supporting information) to obtain pure compound **22e** (0.5 g, 34%). ESI-MS m/z 351 $[M+H]^+$ calc. for $C_{18}H_{23}ClN_2O_3$. This intermediate was used in the next step without further characterization.

4-Chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-(trifluoromethyl)quinoline (22f)

Compound **21** (370 mg, 1 mmol) was dissolved in POCl₃ (30 mL) and stirred at 110 °C for 2 hours. Then, the reaction mixture was concentrated, quenched by ice-water and extracted with EtOAc. The organic phase was dried with Na₂SO₄, filtered and concentrated to give the desired compound **22f** (200 mg, 52%). ESI-MS m/z 389 $[M+H]^+$ calc. for $C_{18}H_{20}ClF_3N_2O_2$. This intermediate was used in the next step without further characterization.

4-Chloro-6-methoxy-2-phenyl-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22g)

To a solution of compound **16** (300 mg, 0.85 mmol) in 1,4-dioxane/H₂O (5:1, 18 mL) was added Na₂CO₃ (180 mg, 1.7 mmol), Pd(PPh₃)₄ (98 mg, 0.085 mmol) and phenylboronic acid (93 mg, 0.76 mmol) and the solution was heated to 110 °C for 4 hours under microwave irradiation. Then, the mixture was concentrated and extracted with EtOAc. The combined organic layers were washed with brine, dried over

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3 anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
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5 purified by preparative TLC to obtain pure compound **22g** (0.1 g, 30%) as pale yellow
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7 solid. ESI-MS *m/z* 397 [M+H]⁺ calc. for C₂₃H₂₅ClN₂O₂. This intermediate was used in
8
9 the next step without further characterization.
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11 12 13 **4-Chloro-6-methoxy-2-(o-tolyl)-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22h)**

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15 To a solution of compound **16** (200 mg, 0.56 mmol) in 1,4-dioxane/H₂O (5:1, 18 mL)
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17 was added K₂CO₃ (78 mg, 0.56 mmol), Pd(PPh₃)₄ (65 mg, 0.056 mmol) and o-
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19 tolylboronic acid (103 mg, 0.76 mmol) and the solution was heated to 110 °C for 4 hours
20
21 under microwave irradiation. Then, the mixture was concentrated and extracted with
22
23 EtOAc. The combined organic layers were washed with brine, dried over anhydrous
24
25 Na₂SO₄, filtered and concentrated to give the crude product which was purified by
26
27 preparative TLC to obtain pure compound **22h** (0.11 g, 48%) as pale yellow solid. ESI-
28
29 MS *m/z* 411 [M+H]⁺ calc. for C₂₄H₂₇ClN₂O₂. This intermediate was used in the next
30
31 step without further characterization.
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37 38 **4-Chloro-6-methoxy-2-(3-pyridyl)-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22i)**

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40 To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was
41
42 added Na₂CO₃ (212 mg, 2 mmol), Pd(PPh₃)₄ (40 mg, 0.03 mmol) and pyridin-3-
43
44 ylboronic acid (123 mg, 1 mmol) and the solution was heated to 110 °C for 2 hours
45
46 under microwave irradiation. Then, the reaction was quenched with water and extracted
47
48 with EtOAc. The organic layer was washed with brine, dried over anhydrous Na₂SO₄,
49
50 filtered and concentrated to give the crude product which was purified by preparative
51
52 HPLC (method 2 described in supporting information) to obtain pure compound **22i**
53
54 (200 mg, 50%) as a yellow solid. ¹H NMR (MeOD, 400 MHz): δ 9.67 (s, 1H), 9.41 (d, *J*

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3 = 8.4 Hz, 1H), 8.93 (d, J = 8.4 Hz, 1H), 8.33 (s, 1H), 8.26-8.22 (m, 1H), 7.63 (s, 1H),
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5 7.60 (s, 1H), 4.45-4.35 (m, 2H), 4.09 (s, 3H), 3.88-3.78 (m, 2H), 3.54-3.48 (m, 2H),
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7 3.22-3.10 (m, 2H), 2.45-2.35 (m, 2H), 2.26-2.15 (m, 2H), 2.11-2.02 (m, 2H). ESI-MS
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9 m/z 398.2 $[M+H]^+$ calc. for $C_{22}H_{24}ClN_3O_2$.

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13 **4-[4-Chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-quinoly]-1H-pyridin-2-one**
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15 **(22j)**

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17 To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was
18
19 added K₂CO₃ (278 mg, 2 mmol), Pd(PPh₃)₄ (50 mg, 0.04 mmol) and 4-(4,4,5,5-
20
21 tetramethyl-1,3,2-dioxaborolan-2-yl)-1H-pyridin-2-one (**Int. 1**) (freshly prepared, 1.25
22
23 mmol in 10 mL of 1,4-dioxane, synthesis described in supporting information) and the
24
25 solution was heated to 120 °C for 4 hours under microwave irradiation. Then, the
26
27 mixture was quenched with water and extracted with EtOAc. The organic layer was
28
29 washed with brine, dried over anhydrous Na₂SO₄ and concentrated to give the crude
30
31 product which was purified by preparative HPLC (method 3 described in supporting
32
33 information) to obtain pure compound **22j** (207 mg, 50%). ¹H NMR (MeOD, 400
34
35 MHz): δ 8.12 (s, 1H), 7.72 (s, 1H), 7.58-7.55 (m, 2H), 7.37 (br s, 2H), 4.43-4.35 (m,
36
37 2H), 4.10 (s, 3H), 3.89-9.78 (m, 2H), 3.55-3.45 (m, 2H), 3.25-3.15 (m, 2H), 2.45-2.35
38
39 (m, 2H), 2.25-2.15 (m, 2H), 2.12-2.02 (m, 2H). ESI-MS m/z 414.2 $[M+H]^+$ calc. for
40
41 $C_{22}H_{24}ClN_3O_3$.

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48 **4-Chloro-2-(3-furyl)-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)quinoline (22k)**
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50 To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was
51
52 added Na₂CO₃ (212 mg, 2 mmol), Pd(PPh₃)₄ (40 mg, 0.03 mmol) and furan-3-ylboronic
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54 acid (159 mg, 1.43 mmol) and the solution was heated to 110 °C for 2 hours under
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3 microwave irradiation. Then, the reaction was quenched with water and extracted with
4 EtOAc. The organic layer was washed with brine, dried over anhydrous Na₂SO₄,
5 filtered and concentrated to give the crude product which was purified by preparative
6 HPLC (method 2 described in supporting information) to obtain pure compound **22k**
7 (120 mg, 31%) as a yellow solid. ¹H NMR (DMSO-*d*₆, 400 MHz): δ 8.51 (s, 1H), 8.00
8 (s, 1H), 7.81 (s, 1H), 7.43-3.37 (m, 2H), 7.15 (s, 1H), 4.33-4.24 (m, 2H), 3.97 (s, 3H),
9 3.68-3.58 (m, 2H), 3.35-3.25 (m, 2H), 3.10-3.00 (m, 2H), 2.25-2.15 (m, 2H), 2.06-1.95
10 (m, 2H), 1.90-1.78 (m, 2H). ESI-MS *m/z* 387.2 [M+H]⁺ calc. for C₂₁H₂₃ClN₂O₃.

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22 **4-Chloro-6-methoxy-2-(5-methyl-2-thienyl)-7-(3-pyrrolidin-1-ylpropoxy)quinoline**
23 **(22l)**

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26 To a solution of compound **16** (200 mg, 0.56 mmol) in 1,4-dioxane/H₂O (5:1, 18 mL)
27 was added K₂CO₃ (116 mg, 0.84 mmol), Pd(PPh₃)₄ (65 mg, 0.056 mmol) and 4,4,5,5-
28 tetramethyl-2-(5-methyl-2-thienyl)-1,3,2-dioxaborolane (126 mg, 0.56 mmol) and the
29 solution was heated to 110 °C for 4 hours under microwave irradiation. Then, the
30 mixture was concentrated and extracted with EtOAc. The combined organic layers were
31 washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
32 crude product which was purified by preparative TLC to obtain pure compound **22l**
33 (0.12 g, 51%) as pale yellow solid. ESI-MS *m/z* 417 [M+H]⁺ calc. for C₂₂H₂₅ClN₂O₂S.
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35 This intermediate was used in the next step without further characterization.
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48 **4-Chloro-6-methoxy-2-(5-methyl-1H-pyrrol-2-yl)-7-(3-pyrrolidin-1-**
49 **ylpropoxy)quinoline (22m)**

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52 To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was
53 added K₂CO₃ (278 mg, 2 mmol), Pd(PPh₃)₄ (50 mg, 0.04 mmol) and 2-methyl-5-
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3 (4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)-1*H*-pyrrole (**Int. 2**, synthesis described in
4 supporting information) (207 mg, 1 mmol) and the solution was heated to 120 °C for 4
5 hours under microwave irradiation. Then, the mixture was quenched with water and
6 extracted with EtOAc. The organic layer was washed with brine, dried over anhydrous
7 Na₂SO₄, filtered and concentrated to give the crude product which was purified by
8 preparative HPLC (method 3 described in supporting information) to obtain pure
9 compound **22m** (150 mg, 38%) as a yellow solid. ¹H NMR (MeOD, 400 MHz): δ 8.13
10 (s, 1H), 7.67 (s, 1H), 7.60 (s, 1H), 7.41 (d, *J* = 4 Hz, 1H), 6.28 (d, *J* = 4 Hz, 1H), 4.48-
11 4.40 (m, 2H), 4.10 (s, 3H), 3.90-3.80 (m, 2H), 3.56-3.48 (m, 2H), 3.26-3.10 (m, 2H),
12 2.50-2.35 (m, 5H), 2.30-2.18 (m, 2H), 2.11-2.02 (m, 2H). ESI-MS *m/z* 400.2 [M+H]⁺
13 calc. for C₂₂H₂₆ClN₃O₂.

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29 **4-Chloro-6-methoxy-2-(2-methylpyrazol-3-yl)-7-(3-pyrrolidin-1-
30 ylpropoxy)quinoline (22n)**

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32 To a solution of compound **16** (270 mg, 0.76 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL)
33 was added Na₂CO₃ (322 mg, 3.04 mmol), Pd(PPh₃)₄ (87.7 mg, 0.07 mmol) and
34 commercially available 1-methyl-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)-1*H*-
35 pyrazole (159 mg, 0.76 mmol). The solution was heated to 110 °C for 2 hours under
36 microwave irradiation. The mixture was quenched with water and extracted with
37 EtOAc. The organic layer was washed with brine, dried over anhydrous Na₂SO₄,
38 filtered and concentrated to give the crude product which was purified by preparative
39 HPLC (method 2 described in supporting information) to obtain pure compound **22n**
40 (120 mg, 40%) as a yellow solid. ESI-MS *m/z* 401 [M+H]⁺ calc. for C₂₁H₂₅ClN₄O₂.
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42 This intermediate was used in the next step without further characterization.
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3 **4-Chloro-6-methoxy-2-(1-methylpyrazol-3-yl)-7-(3-pyrrolidin-1-**
4 **ylpropoxy)quinoline (22o)**
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7 To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was
8 added K₂CO₃ (278 mg, 2 mmol), Pd(PPh₃)₄ (50 mg, 0.05 mmol) and 1-methyl-3-
9 (4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyrazole (207 mg, 1 mmol) and the
10 solution was heated to 120 °C for 4 hours under microwave irradiation. Then, the
11 mixture was quenched with water and extracted with EtOAc. The organic layer was
12 washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
13 crude product which was purified by preparative HPLC (method 3 described in
14 supporting information) to obtain pure compound **22o** (130 mg, 33%) as a yellow solid.
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16 ¹H NMR (MeOD, 400 MHz): δ 8.30 (s, 1H), 7.83 (s, 1H), 7.68 (s, 1H), 7.63 (s, 1H),
17 7.17 (s, 1H), 4.48-4.38 (m, 2H), 4.14-4.03 (m, 6H), 3.85-3.75 (m, 2H), 3.56-3.47 (m,
18 2H), 3.26-3.15 (m, 2H), 2.48-2.35 (m, 2H), 2.30-2.15 (m, 2H), 2.15-2.03 (m, 2H). ESI-
19 MS *m/z* 401.2 [M+H]⁺ calc. for C₂₁H₂₅ClN₄O₂.
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35 **4-Chloro-2-(3,6-dihydro-2H-pyran-4-yl)-6-methoxy-7-(3-pyrrolidin-1-**
36 **ylpropoxy)quinoline (22p)**
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39 To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was
40 added Na₂CO₃ (212 mg, 2 mmol), Pd(PPh₃)₄ (50 mg, 0.04 mmol) and 2-(3,6-dihydro-
41 2H-pyran-4-yl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane (210 mg, 1 mmol) and the
42 solution was heated to 110 °C for 4 hours under microwave irradiation. Then, the
43 reaction was quenched with water and extracted with EtOAc. The organic layer was
44 washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
45 crude product which was purified by preparative HPLC (method 2 described in
46 supporting information) to obtain pure compound **22p** (250 mg, 62%) as a yellow solid.
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¹H NMR (MeOD, 400 MHz): δ 7.92 (s, 1H), 7.59 (s, 1H), 7.57 (s, 1H), 6.90 (s, 1H), 4.43-4.37 (m, 4H), 4.07 (s, 3H), 3.99-3.91 (m, 2H), 3.87-3.77 (m, 2H), 3.51-3.45 (m, 2H), 3.20-3.10 (m, 2H), 2.75-2.70 (m, 2H), 2.55-2.30 (m, 2H), 2.25-2.15 (m, 2H), 2.11-2.03 (m, 2H). ESI-MS *m/z* 403.2 [M+H]⁺ calc. for C₂₂H₂₇ClN₂O₃.

4-Chloro-6-methoxy-2-(5-methyl-2-furyl)-7-(3-pyrrolidin-1-ylpropoxy)quinoline

(22r)

To a mixture of **16** (10.00 g, 28.15 mmol) and 4,4,5,5-tetramethyl-2-(5-methyl-2-furyl)-1,3,2-dioxaborolane (6.44 g, 30.97 mmol) in 1,4-dioxane (100 mL), was added K₂CO₃ (7.78 g, 56.30 mmol) and Pd(PPh₃)₄ (2.00 g, 1.73 mmol) in one portion at 25 °C under N₂ and the mixture was stirred at 25 °C for 10 minutes. Then, the reaction mixture was heated to 120 °C for 12 hours. Then, the mixture was cooled to 25 °C and extracted with EtOAc. The combined organic phase was washed with saturated brine, dried with anhydrous Na₂SO₄, filtered and concentrated in vacuum to give a residue which was purified by column chromatography to afford pure compound **22r** (8.00 g, 71%) as yellow solid. ¹H NMR (DMSO-*d*₆, 400 MHz): δ 7.83 (s, 1H), 7.47 (s, 1H), 7.36 (s, 1H), 7.19 (d, *J* = 3.2 Hz, 1H), 6.31 (d, *J* = 3.2 Hz, 1H), 4.32-4.23 (m, 2H), 3.96 (s, 3H), 3.65-3.55 (m, 2H), 3.36-3.26 (m, 2H), 3.10-3.00 (m, 2H), 2.39 (s, 3H), 2.25-2.15 (m, 2H), 2.06-1.95 (m, 2H), 1.90-1.80 (m, 2H). ESI-MS *m/z* 401.2 [M+H]⁺ calc. for C₂₂H₂₅ClN₂O₃.

2-Cyclohexyl-6-methoxy-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (23a)

To a solution of compound **23b** (47.8 mg, 0.1 mmol) in EtOH (10 mL) was added Pd/C (15 mg) under H₂ (1 atm) and the mixture was stirred at 25 °C for 15 hours. Then, the

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3 reaction mixture was filtered and the filtrate was concentrated to give the desired
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5 compound **23a** (48 mg, 99%) as yellow solid; m.p. 101-102 °C. ¹H NMR (MeOD, 400
6
7 MHz): δ 7.82 (s, 1H), 7.35 (s, 1H), 6.76 (s, 1H), 4.40-4.30 (m, 3H), 4.04 (s, 3H), 3.90-
8
9 3.80 (m, 2H), 3.75-3.66 (m, 2H), 3.55-3.45 (m, 2H), 3.35-3.25 (m, 2H), 3.21-3.12 (m,
10
11 2H), 2.96 (s, 3H), 2.91-2.85 (m, 1H), 2.43-2.31 (m, 4H), 2.22-1.90 (m, 10H), 1.90-1.70
12
13 (m, 3H), 1.58-1.37 (m, 3H). ESI-MS *m/z* 482.5 [M+H]⁺ calc. for C₂₉H₄₄N₄O₂.

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18 **2-(Cyclohexen-1-yl)-6-methoxy-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-**
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20 **ylpropoxy)quinolin-4-amine (23b)**

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22 To a solution of compound **17** (0.33 g, 0.83 mmol) in 1,4-dioxane (15 mL) was added
23
24 Cs₂CO₃ (1.7 g, 5.2 mmol), BINAP (0.154 g, 0.25 mmol), Pd₂(dba)₃ (0.14 g, 0.15 mmol)
25
26 and 1-methylpiperidin-4-amine (0.282 g, 2.48 mmol) and the solution was heated to
27
28 110 °C for 1 hour under microwave irradiation. Then, the reaction mixture was
29
30 concentrated and extracted with EtOAc. The combined organic layers were washed with
31
32 brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product
33
34 which was purified by preparative HPLC (method 1 described in supporting
35
36 information) to obtain pure compound **23b** (80 mg, 20%) as yellow solid; m.p. 189-190
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38 °C. ¹H NMR (MeOD, 400 MHz): δ 7.82 (s, 1H), 7.43 (s, 1H), 6.81 (s, 1H), 6.73 (s, 1H),
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40 4.34-4.25 (m, 3H), 4.05 (s, 3H), 3.85-3.75 (m, 2H), 3.70-3.60 (m, 2H), 3.60-3.40 (m,
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42 2H), 3.30-3.20 (m, 2H), 3.19-3.10 (m, 2H), 2.99 (s, 3H), 2.62-2.55 (m, 2H), 2.40-2.30
43
44 (m, 6H), 2.20-2.10 (m, 6H), 1.91-1.79 (m, 4H). ESI-MS *m/z* 479.6 [M+H]⁺ calc. for
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46 C₂₉H₄₂N₄O₂.

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53 **6-Methoxy-2-(4-methylpiperazin-1-yl)-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-**
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55 **ylpropoxy)quinolin-4-amine (23c)**

To a solution of compound **22c** (0.2 g, 0.48 mmol) in 1,4-dioxane (10 mL) was added Cs₂CO₃ (0.7 g, 2.15 mmol), BINAP (0.059 g, 0.096 mmol), Pd₂(dba)₃ (0.087 g, 0.096 mol) and 1-methylpiperidin-4-amine (0.272 g, 2.39 mmol) and the mixture was heated to 120 °C for one hour under microwave irradiation. Then, the solution was concentrated and extracted with EtOAc. The combined organic layers were washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative HPLC (method 1 described in supporting information) to obtain pure compound **23c** (0.02 g, 9%) as yellow solid; m.p. 107-108 °C. ¹H NMR (MeOD, 400 MHz): δ 7.70 (s, 1H), 7.37 (s, 1H), 6.17 (s, 1H), 4.34-4.25 (m, 3H), 4.25-4.05 (m, 3H), 3.97 (s, 3H), 3.85-3.75 (m, 3H), 3.70-3.60 (m, 2H), 3.60-3.40 (m, 6H), 3.20-3.10 (m, 3H), 2.99 (s, 3H), 2.92 (s, 3H), 2.40-2.30 (m, 4H), 2.25-2.15 (m, 3H), 2.12-2.00 (m, 4H). ESI-MS *m/z* 497.4 [M+H]⁺ calc. for C₂₈H₄₄N₆O₂.

6-Methoxy-2-methyl-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (23d)

To a solution of compound **22d** (100 mg, 0.30 mmol) in DME (5 mL) was added K₃PO₄ (0.11 g, 0.5 mmol), biphenyl-2-yl-dicyclohexyl-phosphane (0.052 g, 0.15 mmol), Pd₂(dba)₃ (0.14 g, 0.015 mmol) and 1-methylpiperidin-4-amine (0.17 g, 1.5 mmol) and the mixture was heated to 110 °C for 3 hours under microwave irradiation. Then, the solution was concentrated and extracted with EtOAc. The combined organic layers were washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative HPLC (method 1 described in supporting information) to obtain pure compound **23d** (25 mg, 20%) as a yellow syrup. ¹H NMR (MeOD, 400 MHz): δ 7.78 (s, 1H), 7.21 (s, 1H), 6.78 (s, 1H), 4.32 (t, *J* = 5.2 Hz, 2H), 4.20-4.17 (m, 1H), 4.00 (s, 3H), 3.82-3.78 (m, 2H), 3.70-3.63 (m, 2H), 3.48 (

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3 t, $J = 6.8$ Hz, 2H), 3.29-3.21 (m, 2H), 3.19-3.11 (m, 2H), 2.95-2.93 (s, 3H), 2.66 (s, 3H),
4
5 2.38-2.32 (m, 4H), 2.21-2.16 (m, 2H), 2.11-2.03 (m, 4H). ESI-MS m/z 413.3 $[M+H]^+$
6
7 calc. for $C_{24}H_{36}N_4O_2$.
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11 **2,6-Dimethoxy-*N*-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-**
12
13 **amine (23e)**
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15 To a solution of compound **22e** (0.28 g, 0.8 mmol) in 1,4-dioxane (15 mL) was added
16 Cs_2CO_3 (1.04 g, 3.2 mmol), BINAP (150 mg, 0.24 mmol), $Pd_2(dba)_3$ (218 mg, 0.24
17 mmol) and 1-methylpiperidin-4-amine (216 mg, 1.9 mmol) and the solution was heated
18 to 110 °C for 1 hour under microwave irradiation. Then, the solution was concentrated
19 and extracted with EtOAc. The combined organic layer was washed with brine, dried
20 over anhydrous Na_2SO_4 , filtered and concentrated to give the crude product which was
21 purified by preparative HPLC (method 1 described in supporting information) to obtain
22 pure compound **23e** (53 mg, 16%) as yellow solid; m.p. 129-130 °C. 1H NMR (MeOD,
23 400 MHz): δ 7.81 (s, 1H), 7.10 (s, 1H), 6.32 (s, 1H), 4.32-4.24 (m, 2H), 4.20 (s, 3H),
24 4.07 (s, 3H), 3.89-3.78 (m, 2H), 3.75-3.63 (m, 2H), 3.53-3.45 (m, 2H), 3.35-3.23 (m,
25 3H), 3.20-3.10 (m, 2H), 2.96 (s, 3H), 2.45-2.30 (m, 4H), 2.25-2.03 (m, 6H). ESI-MS
26 m/z 429.5 $[M+H]^+$ calc. for $C_{24}H_{36}N_4O_3$.
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44 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)-2-**
45 **(trifluoromethyl)quinolin-4-amine (23f)**
46

47 To a solution of compound **22f** (194 mg, 0.5 mmol) in 1,4-dioxane (15 mL) was added
48 Cs_2CO_3 (488 mg, 1.5 mmol), BINAP (94 mg, 0.15 mmol), $Pd_2(dba)_3$ (78 mg, 0.08
49 mmol) and 1-methylpiperidin-4-amine (171 mg, 1.5 mmol) and the mixture was heated
50 to 110 °C for 3 hours under microwave irradiation. Then the solution was concentrated
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3 and extracted with EtOAc. The combined organic layer was washed with brine, dried
4
5 over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
6
7 purified by preparative HPLC (method 1 described in supporting information) to obtain
8
9 pure compound **23f** (52 mg, 22%) as yellow syrup. ¹H NMR (MeOD, 400 MHz): δ 7.92
10
11 (s, 1H), 7.42 (s, 1H), 7.31 (s, 1H), 4.42-3.34 (m, 3H), 4.08 (s, 3H), 3.90-3.80 (m, 2H),
12
13 3.75-3.68 (m, 2H), 3.55-3.45 (m, 2H), 3.35-3.24 (m, 2H), 3.22-3.12 (m, 2H), 2.92 (s,
14
15 3H), 2.45-2.34 (m, 4H), 2.27-2.12 (m, 4H), 2.10-2.02 (m, 2H). ESI-MS *m/z* 467.3
16
17 [M+H]⁺ calc. for C₂₄H₃₃F₃N₄O₂.

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22 **6-methoxy-N-(1-methyl-4-piperidyl)-2-phenyl-7-(3-pyrrolidin-1-**
23
24 **ylpropoxy)quinolin-4-amine (23g)**

25
26 To a solution of compound **22g** (100 mg, 0.25 mmol) in toluene (10 mL) was added *t*-
27
28 BuOK (1.0 M in THF, 0.37 mL, 0.37 mmol), xantphos (24 mg, 0.04 mmol), Pd₂(dba)₃
29
30 (69 mg, 0.075 mmol) and 1-methylpiperidin-4-amine (167 mg, 1.4 mmol) and the
31
32 solution was heated to 130 °C for 5 hours under microwave irradiation. Then, the
33
34 mixture was quenched with water and extracted with EtOAc. The organic layer was
35
36 washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
37
38 crude product which was purified by preparative HPLC (method 1 described in
39
40 supporting information) to obtain pure compound **23g** (18.2 mg, 15%) as a yellow solid;
41
42 m.p. 186-187 °C. ¹H NMR (MeOD, 400 MHz): δ 7.95-7.93 (m, 2H), 7.88 (s, 1H), 7.67-
43
44 7.65 (m, 3H), 7.43 (s, 1H), 7.10 (s, 1H), 4.41-4.32 (m, 3H), 4.06 (s, 3H), 3.85-3.75 (m,
45
46 3H), 3.70-3.62 (m, 2H), 3.52-3.47 (m, 2H), 3.30-3.20 (m, 1H), 3.20-3.10 (m, 2H), 2.93
47
48 (s, 3H), 2.45-2.35 (m, 4H), 2.23-2.02 (m, 6H). ESI-MS *m/z* 475.3 [M+H]⁺ calc. for
49
50 C₂₉H₃₈N₄O₂.

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3 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(*o*-tolyl)-7-(3-pyrrolidin-1-**
4 **ylpropoxy)quinolin-4-amine (23h)**
5

6
7 To a solution of compound **22h** (98 mg, 0.24 mmol) in 1,4-dioxane (5 mL) was added
8
9 BINAP (15 mg, 0.024 mmol), Pd₂(dba)₃ (22 mg, 0.024 mmol), Cs₂CO₃ (157 mg, 0.48
10
11 mmol) and 1-methylpiperidin-4-amine (54 mg, 0.48 mmol) and the solution was heated
12
13 to 130 °C for 5 hours under microwave irradiation. Then, the mixture was quenched
14
15 with water and extracted with EtOAc. The organic layer was washed with brine, dried
16
17 over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
18
19 purified by preparative HPLC (method 1 described in supporting information) to obtain
20
21 pure compound **23h** (16.4 mg, 14%) as a yellow solid; m.p. 98-99 °C. ¹H NMR (MeOD,
22
23 400 MHz): δ 7.90 (s, 1H), 7.51-7.48 (m, 2H), 7.45-7.40 (m, 2H), 7.23 (s, 1H), 6.91 (s,
24
25 1H), 4.35-4.15 (m, 3H), 4.07 (s, 3H), 3.87-3.77 (m, 2H), 3.67-3.58 (m, 2H), 3.49-3.42
26
27 (m, 2H), 3.19-3.14 (m, 4H), 2.89 (s, 3H), 2.42-2.32 (m, 7H), 2.25-2.12 (m, 4H), 2.10-
28
29 2.06 (m, 2H). ESI-MS *m/z* 489.3 [M+H]⁺ calc. for C₃₀H₄₀N₄O₂.
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35 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(3-pyridyl)-7-(3-pyrrolidin-1-**
36 **ylpropoxy)quinolin-4-amine (23i)**
37

38
39 To a solution of compound **22i** (80 mg, 0.2 mmol) in toluene (10 mL) was added *t*-
40
41 BuOK (1.0 M in THF, 0.4 mL, 0.4 mmol), xantphos (9 mg, 0.02 mmol), Pd₂(dba)₃ (30
42
43 mg, 0.03 mmol) and 1-methylpiperidin-4-amine (34 mg, 0.3 mmol) and the solution
44
45 was heated to 130 °C for 3 hours under microwave irradiation. Then, the solution was
46
47 concentrated and extracted with EtOAc. The combined organic layer was washed with
48
49 brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product
50
51 which was purified by preparative HPLC (method 2 described in supporting
52
53 information) to obtain pure compound **23i** (30 mg, 31%) as yellow solid; m.p. 157-158
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2 °C. ¹H NMR (MeOD, 400 MHz): δ 9.18 (br s, 1H), 8.88 (br s, 1H), 8.49-8.47 (m, 1H),
3 7.92 (s, 1H), 7.78 (br s, 1H), 7.45 (s, 1H), 7.21 (s, 1H), 4.44-4.35 (m, 2H), 4.09 (s, 3H),
4 3.90-3.80 (m, 2H), 3.75-3.67 (m, 2H), 3.55-3.45 (m, 2H), 3.38-3.25 (m, 3H), 3.24-3.15
5 (m, 2H), 2.96 (s, 3H), 2.47-2.35 (m, 4H), 2.26-2.05 (m, 6H). ESI-MS *m/z* 476.3 [M+H]⁺
6
7
8
9
10
11 calc. for C₂₈H₃₇N₅O₂.

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16 **4-[6-Methoxy-4-[(1-methyl-4-piperidyl)amino]-7-(3-pyrrolidin-1-ylpropoxy)-2-**
17 **quinolyl]-1*H*-pyridin-2-one (23j)**

18
19 To a solution of compound **22j** (100 mg, 0.24 mmol) in toluene (10 mL) was added *t*-
20 BuOK (1.0 M in THF, 0.5 mL, 0.5 mmol), xantphos (9 mg, 0.02 mmol), Pd₂(dba)₃ (30
21 mg, 0.03 mmol) and 1-methylpiperidin-4-amine (57 mg, 0.5 mmol) and the mixture
22 was heated to 130 °C for 10 hours. Then, the solution was concentrated and extracted
23 with EtOAc. The combined organic layer was washed with brine, dried over anhydrous
24 Na₂SO₄, filtered and concentrated to give the crude product which was purified by
25 preparative HPLC (method 3 described in supporting information) to obtain pure
26 compound **23j** (10 mg, 9%) as yellow solid; m.p. 173-174 °C. ¹H NMR (MeOD, 400
27 MHz): δ 7.91 (s, 1H), 7.73-7.71 (m, 1H), 7.42 (s, 1H), 7.16 (s, 1H), 7.05 (s, 1H), 6.88-
28 6.86 (m, 1H), 4.42-4.33 (m, 3H), 4.09 (s, 3H), 3.90-3.80 (m, 2H), 3.77-3.66 (m, 2H),
29 3.55-3.47 (m, 2H), 3.35-3.13 (m, 4H), 2.96 (s, 3H), 2.45-2.33 (m, 4H), 2.26-2.05 (m,
30 6H). ESI-MS *m/z* 492.3 [M+H]⁺ calc. for C₂₈H₃₇N₅O₃.

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48 **2-(3-Furyl)-6-methoxy-*N*-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-**
49 **ylpropoxy)quinolin-4-amine (23k)**

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51 To a solution of compound **22k** (120 mg, 0.3 mmol) in 1,4-dioxane (10 mL) was added
52 Cs₂CO₃ (325 mg, 1 mmol), BINAP (67 mg, 0.1 mmol), Pd₂(dba)₃ (30 mg, 0.03 mmol)
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3 and 1-methylpiperidin-4-amine (114 mg, 1 mmol) and the mixture was heated to 120 °C
4
5 for 3 hours under microwave irradiation. Then, the solution was concentrated and
6
7 extracted with EtOAc. The combined organic layer was washed with brine, dried over
8
9 anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
10
11 purified by preparative HPLC (method 2 described in supporting information) to obtain
12
13 pure compound **23k** (20 mg, 15%) as yellow solid; m.p. 108-109 °C. ¹H NMR (MeOD,
14
15 400 MHz): δ 8.57 (s, 1H), 7.84-7.81 (m, 2H), 7.45 (s, 1H), 7.25 (s, 1H), 7.07 (s, 1H),
16
17 4.40-4.30 (m, 2H), 4.04 (s, 3H), 3.85-3.75 (m, 2H), 3.72-3.64 (m, 2H), 3.51-3.45 (m,
18
19 2H), 3.30-3.25 (m, 3H), 3.24-3.10 (m, 2H), 2.95 (s, 3H), 2.41-2.34 (m, 4H), 2.23-2.05
20
21 (m, 6H). ESI-MS *m/z* 465.2 [M+H]⁺ calc. for C₂₇H₃₆N₄O₃.

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27 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(5-methyl-2-thienyl)-7-(3-pyrrolidin-1-**
28
29 **ylpropoxy)quinolin-4-amine (23l)**

30
31 To a solution of compound **22l** (100 mg, 0.24 mmol) in 1,4-dioxane (5 mL) was added
32
33 BINAP (15 mg, 0.024 mmol), Pd₂(dba)₃ (22 mg, 0.024 mmol), Cs₂CO₃ (157 mg, 0.48
34
35 mmol) and 1-methylpiperidin-4-amine (54 mg, 0.48 mmol) and the solution was heated
36
37 to 130 °C for 5 hours under microwave irradiation. Then, the mixture was quenched
38
39 with water and extracted with EtOAc. The organic layer was washed with brine, dried
40
41 over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
42
43 purified by preparative HPLC (method 1 described in supporting information) to obtain
44
45 pure compound **23l** (13.4 mg, 11%) as a yellow solid; m.p. 110-111 °C. ¹H NMR
46
47 (MeOD, 400 MHz): δ 7.86-7.83 (m, 2H), 7.44 (s, 1H), 7.04-7.02 (m, 1H), 6.95 (s, 1H),
48
49 4.40-4.25 (m, 3H), 4.04 (s, 3H), 3.85-3.78 (m, 2H), 3.72-3.63 (m, 2H), 3.51-3.44 (m,
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51 2H), 3.21-3.10 (m, 2H), 2.97-2.92 (m, 3H), 2.62 (s, 3H), 2.40-2.32 (m, 4H), 2.21-2.07
52
53 (m, 8H). ESI-MS *m/z* 495.3 [M+H]⁺ calc. for C₂₈H₃₈N₄O₂S.

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5 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(5-methyl-1*H*-pyrrol-2-yl)-7-(3-pyrrolidin-**
6
7 **1-ylpropoxy)quinolin-4-amine (23m)**

8
9 To a solution of compound **22m** (100 mg, 0.25 mmol) in toluene (10 mL) was added *t*-
10
11 BuOK (1.0 M in THF, 0.5 mL, 0.5 mmol), xantphos (9 mg, 0.02 mmol), Pd₂(dba)₃ (30
12
13 mg, 0.03 mmol) and 1-methylpiperidin-4-amine (57 mg, 0.5 mmol) and the solution
14
15 was heated to 130 °C for 3 hours under microwave irradiation. Then, the solution was
16
17 concentrated and extracted with EtOAc. The combined organic layer was washed with
18
19 brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product
20
21 which was purified by preparative HPLC (method 3 described in supporting
22
23 information) to obtain pure compound **23m** (15 mg, 13%) as yellow syrup. ¹H NMR
24
25 (MeOD, 400 MHz): δ 7.78 (s, 1H), 7.42 (br s, 2H), 7.18 (br s, 1H), 6.98 (s, 1H), 6.14 (s,
26
27 1H), 4.40-4.30 (m, 2H), 4.10-3.97 (m, 4H), 3.51-3.40 (m, 8H), 2.80-2.55 (m, 5H), 2.50
28
29 (s, 3H), 2.48-2.30 (m, 2H), 2.29-2.21 (m, 2H), 2.20-1.95 (m, 6H). ESI-MS *m/z* 478.3
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31 [M+H]⁺ calc. for C₂₈H₃₉N₅O₂.

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38 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(2-methylpyrazol-3-yl)-7-(3-pyrrolidin-1-**
39
40 **ylpropoxy)quinolin-4-amine (23n)**

41
42 To a solution of compound **22n** (95 mg, 0.24 mmol) in 1,4-dioxane (5 mL) was added
43
44 Cs₂CO₃ (406 mg, 1.25 mmol), BINAP (34 mg, 0.05 mmol), Pd₂(dba)₃ (27 mg, 0.03
45
46 mmol), and 1-methylpiperidin-4-amine (143 mg, 1.25 mmol) and the solution was
47
48 heated to 120 °C for 3 hours under microwave irradiation. Then, the solution was
49
50 concentrated and extracted with EtOAc. The combined organic layer was washed with
51
52 brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product
53
54 which was purified by preparative HPLC (method 2 described in supporting
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3 information) to obtain pure compound **23n** (40 mg, 35%) as yellow solid; m.p. 80-81
4 °C. ¹H NMR (MeOD, 400 MHz): δ 7.89 (s, 1H), 7.68 (d, *J* = 2.4 Hz, 1H), 7.32 (s, 1H),
5 7.05 (s, 1H), 6.83 (d, *J* = 2.4 Hz, 1H), 4.40-4.30 (m, 2H), 4.06 (s, 3H), 4.03 (s, 3H),
6 3.85-3.77 (m, 2H), 3.70-3.62 (m, 2H), 3.51-3.48 (m, 2H), 3.32-3.10 (m, 5H), 2.92 (s,
7 3H), 2.44-2.30 (m, 4H), 2.25-2.02 (m, 6H). ESI-MS *m/z* 479.2 [M+H]⁺ calc. for
8 C₂₇H₃₈N₆O₂.
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18 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(1-methylpyrazol-3-yl)-7-(3-pyrrolidin-1-**
19 **ylpropoxy)quinolin-4-amine (23o)**
20
21

22 To a solution of compound **22o** (100 mg, 0.25 mmol) in toluene (10 mL) was added *t*-
23 BuOK (1.0 M in THF, 0.5 mL, 0.5 mmol), xantphos (9 mg, 0.02 mmol), Pd₂(dba)₃ (30
24 mg, 0.03 mmol) and 1-methylpiperidin-4-amine (57 mg, 0.5 mmol) and the solution
25 was heated to 130 °C for 3 hours under microwave irradiation. Then, the solution was
26 concentrated and extracted with EtOAc. The combined organic layer was washed with
27 brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product
28 which was purified by preparative HPLC (method 3 described in supporting
29 information) to obtain pure compound **23o** (25 mg, 21%) as yellow solid; m.p. 139-140
30 °C. ¹H NMR (MeOD, 400 MHz): δ 7.84 (br s, 2H), 7.55 (s, 1H), 7.25-7.21 (m, 2H),
31 4.40-4.30 (m, 3H), 4.07 (s, 3H), 4.05 (s, 3H), 3.86-3.76 (m, 2H), 3.75-3.65 (m, 2H),
32 3.53-3.46 (m, 2H), 3.20-3.08 (m, 4H), 2.95 (s, 3H), 2.45-2.30 (m, 4H), 2.25-2.00 (m,
33 6H). ESI-MS *m/z* 479.3 [M+H]⁺ calc. for C₂₇H₃₈N₆O₂.
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50 **2-(3,6-Dihydro-2*H*-pyran-4-yl)-6-methoxy-*N*-(1-methyl-4-piperidyl)-7-(3-**
51 **pyrrolidin-1-ylpropoxy)quinolin-4-amine (23p)**
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To a solution of compound **22p** (200 mg, 0.5 mmol) in 1,4-dioxane (10 mL) was added Cs₂CO₃ (500 mg, 1.5 mmol), BINAP (67.5 mg, 0.1 mmol), Pd₂(dba)₃ (40 mg, 0.04 mmol) and 1-methylpiperidin-4-amine (114 mg, 1 mmol) and the solution was heated to 120 °C for 3 hours under microwave irradiation. Then, the solution was concentrated and extracted with EtOAc. The combined organic layer was washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative HPLC (method 2 described in supporting information) to obtain pure compound **23p** (100 mg, 42%) as yellow solid; m.p. 182-183 °C. ¹H NMR (MeOD, 400 MHz): δ 7.84 (s, 1H), 7.43 (s, 1H), 6.86 (s, 1H), 6.82 (s, 1H), 4.45-4.20 (m, 5H), 4.03 (s, 3H), 4.02-3.97 (m, 2H), 3.85-3.78 (m, 2H), 3.74-3.68 (m, 2H), 3.53-3.48 (m, 2H), 3.25-3.10 (m, 4H), 2.97 (s, 3H), 2.75-2.70 (m, 2H), 2.45-2.33 (m, 4H), 2.30-2.05 (m, 6H). ESI-MS *m/z* 481.3 [M+H]⁺ calc. for C₂₈H₄₀N₄O₃.

6-Methoxy-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)-2-tetrahydropyran-4-yl-quinolin-4-amine (23q)

To a solution of compound **23p** (48 mg, 0.1 mmol) in MeOH (15 mL) was added Pd/C (10 mg) under H₂ (1 atm) and the solution was heated to 35 °C for 4 hours. Then, the mixture was filtrated and the filtrate was concentrated to give compound **23q** (30 mg, 62%) as a yellow solid; m.p. 120-121 °C. ¹H NMR (MeOD, 400 MHz): δ 7.84 (s, 1H), 7.33 (s, 1H), 6.80 (s, 1H), 4.45-4.10 (m, 5H), 4.03 (s, 3H), 3.90-3.78 (m, 2H), 3.75-3.60 (m, 4H), 3.53-3.48 (m, 2H), 3.30-3.15 (m, 5H), 2.97 (s, 3H), 2.41-2.33 (m, 4H), 2.28-1.90 (m, 10H). ESI-MS *m/z* 483.3 [M+H]⁺ calc. for C₂₈H₄₂N₄O₃.

Ethyl 3-(3-benzyloxy-4-methoxy-anilino)-3-oxo-propanoate (25)

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3 To a mixture of commercially available 3-benzyloxy-4-methoxy-aniline (**24**) (35 g,
4 0.153 mol) and Et₃N (30.87 g, 0.306 mol) in CH₂Cl₂ (1 L) was added dropwise ethyl 3-
5 chloro-3-oxo-propanoate (25.245 g, 0.168 mol) at 0 °C and the mixture was stirred at
6 room temperature for 12 hours. Then, the reaction mixture was poured into water (2 L)
7 and extracted with CH₂Cl₂. The combined the organic phase was dried over anhydrous
8 Na₂SO₄, filtered and concentrated to afford compound **25** (40 g, 76%). ESI-MS *m/z*
9 344.2 [M+H]⁺ calc. for C₁₉H₂₁NO₅. This intermediate was used in the next step without
10 further characterization.
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22 **3-(3-Benzyloxy-4-methoxy-anilino)-3-oxo-propanoic acid (26)**

23
24 To a mixture of **25** (20.40 g, 59.41 mmol) in THF/MeOH/H₂O (3:3:2, 267 mL) was
25 added LiOH·H₂O (3.74 g, 89.12 mmol) in one portion at 25 °C and the mixture was
26 stirred at room temperature for 16 hours. Then, the mixture was concentrated by rotary
27 evaporation under vacuum at 45 °C. The residue was poured into ice-water (200 mL)
28 and stirred for 10 minutes. The resulting slurry was filtered and the filter cake was
29 dried under vacuum to afford compound **26** (19.30 g, 99% crude) as a white solid. ESI-
30 MS *m/z* 316.2 [M+H]⁺ calc. for C₁₇H₁₇NO₅. This intermediate was used in the next step
31 without further characterization.
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44 **7-Benzyloxy-2,4-dichloro-6-methoxy-quinoline (27)**

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46 Compound **26** (7.00 g, 22.20 mmol) was suspended in POCl₃ (68.08 g, 443.99 mmol) in
47 a 500 mL single-necked round bottom flask and the mixture was stirred at 90 °C for 2
48 hours under N₂. Then, the reaction mixture was cooled to 25 °C and concentrated to
49 remove POCl₃. The residue was further purified by silica gel column chromatography to
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3 obtain pure compound **27** (2.50 g, 34%). ¹H NMR (CDCl₃, 400 MHz): δ 7.49-7.36 (m,
4 8H), 5.30 (s, 2H), 4.06 (s, 3H). ESI-MS *m/z* 334.2 [M+H]⁺ calc. for C₁₇H₁₃Cl₂NO₂.

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9 **7-Benzyloxy-4-chloro-6-methoxy-2-(5-methyl-2-furyl)quinoline (28)**

10
11 Compound **27** (900 mg, 2.69 mmol), 4,4,5,5-tetramethyl-2-(5-methyl-2-furyl)-1,3,2-
12 dioxaborolane (588 mg, 2.83 mmol), K₂CO₃ (558 mg, 4.04 mmol) and Pd(PPh₃)₄ (311
13 mg, 0.26 mmol) were dissolved in 1,4-dioxane (10 mL) and heated to 100 °C for 16
14 hours under N₂. Then, the reaction mixture was poured into H₂O and was extracted with
15 EtOAc. The organic phase was washed with saturated brine, dried over anhydrous
16 Na₂SO₄, filtered and concentrated in vacuum to give a residue, which was purified by
17 column chromatography to afford pure compound **28** (500 mg, 49%). ESI-MS *m/z*
18 380.1 [M+H]⁺ calc. for C₂₂H₁₈ClNO₃. This intermediate was used in the next step
19 without further characterization.
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33 **7-Benzyloxy-6-methoxy-2-(5-methyl-2-furyl)-N-(1-methyl-4-piperidyl)quinolin-4-**
34 **amine (29)**

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37 Compound **28** (500 mg, 1.32 mmol), 1-methylpiperidin-4-amine (301 mg, 2.63 mmol),
38 Pd₂(dba)₃ (121 mg, 0.13 mmol), BINAP (82 mg, .013 mmol) and Cs₂CO₃ (858 mg, 2.63
39 mmol) were dissolved in 1,4-dioxane (10 mL) and heated at 110 °C for 16 hours under
40 N₂. Then, the reaction mixture was purified by column chromatography to afford pure
41 compound **29** (400 mg, 66%). ESI-MS *m/z* 458.2 [M+H]⁺ calc. for C₂₈H₃₁N₃O₃. This
42 intermediate was used in the next step without further characterization.
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52 **6-Methoxy-4-[(1-methyl-4-piperidyl)amino]-2-(5-methyltetrahydrofuran-2-**
53 **yl)quinolin-7-ol (30)**
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3 A mixture of **29** (400 mg, 0.87 mmol) and Pd/C (100 mg) in MeOH (20 mL) was stirred
4
5 at 50 °C under H₂ (50 Psi) for 16 hours. Then, catalyst was removed by filtration and
6
7 the filtrate was concentrated to dryness to give compound **30** (300 mg, 93%) as a yellow
8
9 solid. ¹H NMR (MeOD, 400 MHz): δ 7.83 (s, 1H), 7.31 (s, 1H), 6.81 (s, 1H), 5.11-5.07
10
11 (m, 1H), 4.32-4.15 (m, 2H), 4.06 (s, 3H), 3.65-3.54 (m, 2H), 3.35-3.20 (m, 2H), 2.89 (s,
12
13 3H), 2.79-2.52 (m, 1H), 2.35-2.10 (m, 5H), 2.06-1.95 (m, 1H), 1.70-1.60 (m, 1H), 1.43
14
15 (d, *J* = 6 Hz, 3H). ESI-MS *m/z* 372.3 [M+H]⁺ calc. for C₂₁H₂₉N₃O₃.

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20 **6-Methoxy-*N*-(1-methyl-4-piperidyl)-2-(5-methyltetrahydrofuran-2-yl)-7-(3-**
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22 **pyrrolidin-1-ylpropoxy)quinolin-4-amine (31)**

23
24 A mixture of **30** (200 mg, 0.53 mmol), 1-(3-chloropropyl)pyrrolidine (95 mg, 0.64
25
26 mmol) and Cs₂CO₃ (351 mg, 1.08 mmol) in DMF (5 mL) was degassed and purged with
27
28 N₂ for 3 times. Then, the mixture was stirred at 100 °C for 16 hours under N₂. Then, the
29
30 mixture was concentrated to give a residue which was purified by preparative HPLC
31
32 (method 1 described in supporting information) to obtain pure compound **31** (50 mg,
33
34 19%) as a yellow syrup (racemic mixture). ¹H NMR (MeOD, 400 MHz): δ 7.82 (s, 1H),
35
36 7.50 (s, 1H), 6.84 (s, 1H), 5.13-5.05 (m, 1H), 4.37-4.12 (m, 4H), 4.03 (s, 3H), 3.86-3.76
37
38 (m, 2H), 3.71-3.65 (m, 2H), 3.52-3.44 (m, 2H), 3.31-3.20 (m, 2H), 3.18-3.10 (m, 2H),
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40 2.94 (s, 3H), 2.63-2.56 (m, 1H), 2.39-2.30 (m, 4H), 2.25-2.13 (m, 4H), 2.08-2.00 (m,
41
42 4H), 1.70-1.60 (m, 1H), 1.45 (d, *J* = 6.4 Hz, 3H). ESI-MS *m/z* 483.4 [M+H]⁺ calc. for
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44 C₂₈H₄₂N₄O₃.

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50 **5-[4-Chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-quinolyl]furan-2-**
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52 **carbaldehyde (32)**

To a solution of compound **16** (0.2 g, 0.56 mmol) in 1,4-dioxane/H₂O (5:1, 12 mL) was added Na₂CO₃ (65 mg, 0.6 mmol), Pd(PPh₃)₄ (65 mg, 0.056 mmol) and (5-formylfuran-2-yl)boronic acid (69 mg, 0.50 mmol) and the solution was heated to 110 °C for 2 hours under microwave irradiation. Then, the mixture was concentrated and extracted with EtOAc. The combined organic layers were washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative TLC to obtain pure compound **32** (100 mg, 43%) as pale yellow solid. ESI-MS *m/z* 415 [M+H]⁺ calc. for C₂₂H₂₃ClN₂O₄. This intermediate was used in the next step without further characterization.

4-Chloro-2-(5-ethyl-2-furyl)-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)quinoline

(33a)

To a solution of compound **16** (8.80 g, 24.77 mmol) in 1,4-dioxane/H₂O (10:1, 110 mL) were successively added 2-(5-ethyl-2-furyl)-4,4,5,5-tetramethyl-1,3,2-dioxaborolane (5.78 g, 26.01 mmol), Pd(PPh₃)₄ (2.86 g, 2.48 mmol) and K₂CO₃ (8.56 g, 61.93 mmol) and the resulting mixture was stirred at 80 °C for 12 hours under N₂. Then, the mixture was diluted with water and extracted with EtOAc. The combined organic phase was washed with brine, dried over anhydrous Na₂SO₄, filtered, concentrated and purified by silica gel column chromatography to obtain pure compound **33a** (9.36 g, 91%) as a yellow solid. ESI-MS *m/z* 415.2 [M+H]⁺ calc. for C₂₃H₂₇ClN₂O₃. This intermediate was used in the next step without further characterization.

***N*-[[5-[4-chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-quinolyl]-2-furyl]methyl]ethanamine (33b)**

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3 To a solution of compound **32** (180 mg, 0.43 mmol) in MeOH (5 mL) was added
4 ethylamine hydrochloride (105 mg, 1.30 mmol) and the solution was stirred at room
5 temperature for 90 minutes. Then, NaBH₃CN (135 mg, 2.15 mmol) was added to the
6 solution and the mixture was stirred at room temperature for 12 hours. Then, the
7 reaction was quenched with water and extracted with EtOAc. The organic layer was
8 washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
9 crude product which was purified by preparative HPLC (method 1 described in
10 supporting information) to obtain pure compound **33b** (80 mg, 42%) as a yellow solid.
11 ESI-MS *m/z* 444.2 [M+H]⁺ calc. for C₂₄H₃₀ClN₃O₃. This intermediate was used in the
12 next step without further characterization.
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26 **1-[5-[4-Chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-quinolyl]-2-furyl]-*N,N*-**
27 **dimethyl-methanamine (33c)**
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29
30 To a solution of compound **32** (180 mg, 0.43 mmol) in MeOH (5 mL) was added
31 dimethylamine hydrochloride (105 mg, 1.30 mmol) and the solution was stirred at room
32 temperature for 90 minutes. Then, NaBH₃CN (135 mg, 2.15 mmol) was added and the
33 mixture was stirred at room temperature for 12 hours. Then, the reaction was quenched
34 with water and extracted with EtOAc. The organic layer was washed with brine, dried
35 over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
36 purified by preparative HPLC (method 1 described in supporting information) to obtain
37 pure compound **33c** (100 mg, 52%) as a yellow solid. ESI-MS *m/z* 444.2 [M+H]⁺ calc.
38 for C₂₄H₃₀ClN₃O₃. This intermediate was used in the next step without further
39 characterization.
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3 **[5-[4-Chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-quinolyl]-2-furyl]methanol**
4
5 **(33d)**

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7 To a solution of compound **32** (200 mg, 0.48 mmol) in MeOH (5 mL) was added
8
9 NaBH₄ (91.2 mg, 2.4 mmol) and the solution was stirred at room temperature for 2
10
11 hours. Then, the mixture was concentrated to give the crude product which was purified
12
13 by preparative TLC to obtain pure compound **33d** (0.1 g, 50%) as pale yellow solid.
14
15 ESI-MS *m/z* 417 [M+H]⁺ calc. for C₂₂H₂₅ClN₂O₄. This intermediate was used in the
16
17 next step without further characterization.
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22 **5-[4-Chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-2-quinolyl]furan-2-**
23
24 **carbonitrile (33e)**

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26 To a solution of compound **32** (100 mg, 0.24 mmol) in MeOH/CH₂Cl₂ (1:4, 10 mL) was
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28 added phenylphosphonic dichloride (94 mg, 0.48 mmol), pyridine (76 mg, 0.96 mmol)
29
30 and NH₂OH·HCl (17 mg, 0.24 mmol) and the solution was stirred at room temperature
31
32 for 15 hours. Then, NaBH₃CN (135 mg, 2.15 mol) was added to the solution and the
33
34 mixture was stirred at room temperature for 12 hours. Then, the mixture was
35
36 concentrated and extracted with EtOAc. The organic layer was washed with aqueous
37
38 NaHCO₃ and brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
39
40 crude product which was purified by column chromatography to obtain pure compound
41
42 **33e** (50 mg, 51%) as a yellow solid. ESI-MS *m/z* 412 [M+H]⁺ calc. for C₂₂H₂₂ClN₃O₃.
43
44 This intermediate was used in the next step without further characterization.
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50 **4-Chloro-6-methoxy-2-(5-phenyl-2-furyl)-7-(3-pyrrolidin-1-ylpropoxy)quinoline**
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52 **(33f)**
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To a solution of compound **16** (354 mg, 1 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL) was added K₂CO₃ (278 mg, 2 mmol), Pd(PPh₃)₄ (50 mg, 0.05 mmol) and 4,4,5,5-tetramethyl-2-(5-phenyl-2-furyl)-1,3,2-dioxaborolane (**Int. 3**, synthesis described in supporting information) (272 mg, 1 mmol) and the solution was heated to 120 °C for 4 hours under microwave irradiation. Then, the mixture was quenched with water and extracted with EtOAc. The organic layer was washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative HPLC (method 3 described in supporting information) to obtain pure compound **33f** (150 mg, 33%) as a yellow solid. ¹H NMR (MeOD, 400 MHz): δ 8.09 (s, 1H), 7.95-7.92 (m, 2H), 7.48 (m, 6H), 7.07 (s, 1H), 4.38-4.28 (m, 2H), 4.06 (s, 3H), 3.86-3.77 (m, 2H), 3.55-3.45 (m, 2H), 3.21-3.10 (m, 2H), 2.45-2.35 (m, 2H), 2.30-2.15 (m, 2H), 2.11-2.01 (m, 2H). ESI-MS *m/z* 463.2 [M+H]⁺ calc. for C₂₇H₂₇ClN₂O₃.

2-(5-Ethyl-2-furyl)-6-methoxy-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (34a)

To a solution of compound **33a** (207 mg, 0.5 mmol) in 1,4-dioxane (15 mL) was added Cs₂CO₃ (325 mg, 1 mmol), BINAP (34 mg, 0.05 mmol), Pd₂(dba)₃ (30 mg, 0.03 mmol) and 1-methylpiperidin-4-amine (114 mg, 1.0 mmol) and the solution was heated to 130 °C for 4 hours under microwave irradiation. Then, the solution was concentrated and extracted with EtOAc. The combined organic layer was washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative HPLC (method 2 described in supporting information) to obtain pure compound **34a** (35 mg, 14%) as yellow solid; m.p. 133-134 °C. ¹H NMR (MeOD, 400 MHz): δ 7.79 (s, 1H), 7.57 (d, *J* = 3.2 Hz, 1H), 7.49 (s, 1H), 7.08 (s, 1H), 6.46 (d, *J* = 3.2 Hz, 1H), 4.38-4.25 (m, 3H), 4.05 (s, 3H), 3.87-3.77 (m, 2H), 3.75-3.65 (m, 2H),

3.51-3.45 (m, 2H), 3.25-3.10 (m, 4H), 2.94 (s, 3H), 2.90-2.80 (m, 2H), 2.45-2.30 (m, 4H), 2.26-2.02 (m, 6H), 1.40-1.32 (m, 3H). ESI-MS m/z 493.3 $[M+H]^+$ calc. for $C_{29}H_{40}N_4O_3$.

2-[5-(ethylaminomethyl)-2-furyl]-6-methoxy-*N*-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (34b)

To a solution of compound **33b** (70 mg, 0.16 mmol) in toluene (5 mL) was added *t*-BuOK (1.0 M in THF, 0.24 mL, 0.24 mmol), xantphos (15 mg, 0.026 mmol), $Pd_2(dba)_3$ (44 mg, 0.048 mmol) and 1-methylpiperidin-4-amine (106 mg, 0.9 mmol) and the solution was heated to 130 °C for 2 hours under microwave irradiation. Then, the mixture was quenched with water and extracted with EtOAc. The organic layer was washed with brine, dried over anhydrous Na_2SO_4 , filtered and concentrated to give the crude product which was purified by preparative HPLC (method 1 described in supporting information) to obtain pure compound **34b** (9.7 mg, 11%) as a yellow syrup. 1H NMR (MeOD, 400 MHz): δ 7.84 (s, 1H), 7.72-7.71 (m, 1H), 7.66 (s, 1H), 7.22 (s, 1H), 6.97-6.96 (m, 1H), 4.46 (s, 2H), 4.40-4.30 (m, 3H), 4.05 (s, 3H), 3.86-3.76 (m, 2H), 3.74-3.65 (m, 2H), 3.52-3.44 (m, 2H), 3.32-3.23 (m, 2H), 3.21-3.12 (m, 4H), 2.96 (s, 3H), 2.45-2.33 (m, 4H), 2.25-2.15 (m, 4H), 2.15-2.00 (m, 2H), 1.40-1.34 (m, 3H). ESI-MS m/z 522.3 $[M+H]^+$ calc. for $C_{30}H_{43}N_5O_3$.

2-[5-[(Dimethylamino)methyl]-2-furyl]-6-methoxy-*N*-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (34c)

To a solution of compound **33c** (80 mg, 0.18 mmol) in toluene (5 mL) was added *t*-BuOK (1.0 M in THF, 0.27 mL, 0.27 mmol), xantphos (17 mg, 0.03 mmol), $Pd_2(dba)_3$ (49 mg, 0.054 mmol) and 1-methylpiperidin-4-amine (119 mg, 1 mmol) and the

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3 solution was heated to 130 °C for 2 hours under microwave irradiation. Then, the
4
5 mixture was quenched with water and extracted with EtOAc. The organic layer was
6
7 washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the
8
9 crude product which was purified by preparative HPLC (method 1 described in
10
11 supporting information) to obtain pure compound **34c** (20.7 mg, 22%) as a yellow
12
13 syrup. ¹H NMR (MeOD, 400 MHz): δ 7.84 (s, 1H), 7.74 (s, 1H), 7.65 (s, 1H), 7.23 (s,
14
15 1H), 7.07-7.06 (m, 1H), 4.58 (s, 2H), 4.40-4.30 (m, 3H), 4.05 (s, 3H), 3.85-3.75 (m,
16
17 2H), 3.75-3.60 (m, 2H), 3.50-3.42 (m, 2H), 3.36-3.28 (m, 2H), 3.19-3.08 (m, 2H), 2.97-
18
19 2.95 (m, 6H), 2.94 (s, 3H), 2.43-2.35 (m, 4H), 2.25-2.15 (m, 4H), 2.15-2.02 (m, 2H).
20
21 ESI-MS *m/z* 522.3 [M+H]⁺ calc. for C₃₀H₄₃N₅O₃.
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27 **[5-[6-Methoxy-4-[(1-methyl-4-piperidyl)amino]-7-(3-pyrrolidin-1-ylpropoxy)-2-**
28
29 **quinolyl]-2-furyl]methanol (34d)**

30
31 To a solution of compound **33d** (90 mg, 0.21 mmol) in 1,4-dioxane (4 mL) was added
32
33 Cs₂CO₃ (0.21 g, 0.65 mmol), BINAP (27 mg, 0.043 mmol), Pd₂(dba)₃ (59 mg, 0.065
34
35 mmol) and 1-methylpiperidin-4-amine (86 mg, 0.75 mmol) and the solution was heated
36
37 to 120 °C for 5 hours under microwave irradiation. Then, the mixture was quenched
38
39 with water and extracted with EtOAc. The organic layer was washed with brine, dried
40
41 over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was
42
43 purified by preparative HPLC (method 1 described in supporting information) to obtain
44
45 pure compound **34d** (10 mg, 10%) as a yellow solid; m.p. 104-105 °C. ¹H NMR
46
47 (MeOD, 400 MHz): δ 7.83 (s, 1H), 7.64-7.63 (m, 1H), 7.49 (s, 1H), 7.17 (s, 1H), 6.68-
48
49 6.67 (m, 1H), 4.71 (s, 2H), 4.43-4.28 (m, 3H), 4.04 (s, 3H), 3.88-3.78 (m, 2H), 3.75-
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51 3.65 (m, 2H), 3.52-3.43 (m, 2H), 3.30-3.25 (m, 2H), 3.20-3.08 (m, 2H), 2.95 (s, 3H),
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2.43-2.34 (m, 4H), 2.25-2.16 (m, 4H), 2.15-2.02 (m, 2H). ESI-MS m/z 495.3 $[M+H]^+$
calc. for $C_{28}H_{38}N_4O_4$.

5-[6-Methoxy-4-[(1-methyl-4-piperidyl)amino]-7-(3-pyrrolidin-1-ylpropoxy)-2-quinolyl]furan-2-carbonitrile (34e)

To a solution of compound **33e** (80 mg, 0.19 mmol) in 1,4-dioxane (5 mL) was added Cs_2CO_3 (127 mg, 0.39 mmol), BINAP (12 mg, 0.019 mmol), $Pd_2(dba)_3$ (18 mg, 0.019 mmol) and 1-methylpiperidin-4-amine (44 mg, 0.38 mmol) and the solution was heated to 130 °C for 2 hours under microwave irradiation. Then, the mixture was quenched with water and extracted with EtOAc. The organic layer was washed with brine, dried over anhydrous Na_2SO_4 , filtered and concentrated to give the crude product which was purified by preparative HPLC (method 1 described in supporting information) to obtain pure compound **34e** (11.4 mg, 12%) as a yellow solid; m.p. 172-174 °C. 1H NMR (MeOD, 400 MHz): δ 7.86 (s, 1H), 7.83-7.82 (m, 1H), 7.65-7.64 (m, 1H), 7.53 (s, 1H), 7.31 (s, 1H), 4.42-4.33 (m, 3H), 4.05 (s, 3H), 3.90-3.80 (m, 2H), 3.75-3.65 (m, 2H), 3.52-3.45 (m, 2H), 3.32-3.30 (m, 2H), 3.20-3.10 (m, 2H), 2.96 (s, 3H), 2.45-2.34 (m, 4H), 2.25-2.12 (m, 4H), 2.11-2.01 (m, 2H). ESI-MS m/z 490.2 $[M+H]^+$ calc. for $C_{28}H_{35}N_5O_3$.

6-Methoxy-N-(1-methyl-4-piperidyl)-2-(5-phenyl-2-furyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (34f)

To a solution of compound **33f** (100 mg, 0.22 mmol) in toluene (10 mL) was added *t*-BuOK (1.0 M in THF, 0.5 mL, 0.5 mmol), xantphos (9 mg, 0.02 mmol), $Pd_2(dba)_3$ (30 mg, 0.03 mmol) and 1-methylpiperidin-4-amine (57 mg, 0.5 mmol) and the mixture was heated to 130 °C for 3 hours under microwave irradiation. Then, the solution was

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2
3 concentrated and extracted with EtOAc. The combined organic layer was washed with
4
5 brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product
6
7 which was purified by preparative HPLC (method 3 described in supporting
8
9 information) to obtain pure compound **34f** (21 mg, 18%) as yellow solid; m.p. 207-208
10
11 °C. ¹H NMR (MeOD, 400 MHz): δ 8.04-8.02 (m, 2H), 7.86-7.79 (m, 2H), 7.60-7.40 (m,
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13 4H), 7.23-7.20 (m, 2H), 4.45-4.35 (m, 3H), 4.08 (s, 3H), 3.90-3.85 (m, 2H), 3.78-3.70
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15 (m, 2H), 3.56-3.48 (m, 2H), 3.25-3.15 (m, 4H), 3.00 (s, 3H), 2.47-2.38 (m, 4H), 2.30-
16
17 2.05 (m, 6H). ESI-MS *m/z* 541.3 [M+H]⁺ calc. for C₃₃H₄₀N₄O₃.

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22 **2-(Benzofuran-2-yl)-4-chloro-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)quinoline**
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24 **(35)**

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26 To a solution of compound **16** (280 mg, 0.79 mmol) in 1,4-dioxane/H₂O (15:1, 16 mL)
27
28 was added Na₂CO₃ (335 mg, 3.16 mmol), Pd(PPh₃)₄ (92 mg, 0.079 mmol) and
29
30 benzofuran-2-ylboronic acid (128 mg, 0.79 mmol) and the solution was heated to 110 °C
31
32 for 1 hour under microwave irradiation. Then, the mixture was quenched with water and
33
34 extracted with EtOAc. The organic layer was washed with brine, dried over anhydrous
35
36 Na₂SO₄, filtered and concentrated to give the crude product which was purified by
37
38 preparative HPLC (method 2 described in supporting information) to obtain pure
39
40 compound **35** (250 mg, 73%) as a yellow solid. ¹H NMR (DMSO-*d*₆, 400 MHz): δ 9.52-
41
42 9.45 (m, 1H), 8.18 (s, 1H), 7.80-7.65 (m, 2H), 7.57 (s, 1H), 7.45-7.35 (m, 2H), 7.32-
43
44 7.25 (m, 1H), 4.35-4.28 (m, 2H), 3.99 (s, 3H), 3.38-3.30 (m, 2H), 3.10-3.02 (m, 2H),
45
46 2.27-2.18 (m, 2H), 2.09-1.98 (m, 2H), 1.90-1.80 (m, 2H), 1.60-1.50 (m, 2H).
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52 **2-(Benzofuran-2-yl)-6-methoxy-N-(1-methyl-4-piperidyl)-7-(3-pyrrolidin-1-**
53
54 **ylpropoxy)quinolin-4-amine (36)**

To a solution of compound **35** (210 mg, 0.5 mmol) in 1,4-dioxane (10 mL) was added Cs₂CO₃ (815 mg, 2.5 mmol), BINAP (62.5 mg, 0.1 mmol), Pd₂(dba)₃ (20 mg, 0.02 mmol) and 1-methylpiperidin-4-amine (285 mg, 2.5 mmol) and the mixture was heated to 120 °C for 3 hours under microwave irradiation. Then, the solution was concentrated and extracted with EtOAc. The combined organic layer was washed with brine, dried over anhydrous Na₂SO₄, filtered and concentrated to give the crude product which was purified by preparative HPLC (method 2 described in supporting information) to obtain pure compound **36** (60 mg, 23%) as yellow solid; m.p. 191-192 °C. ¹H NMR (MeOD, 400 MHz): δ 8.09 (s, 1H), 7.87 (s, 1H), 7.81 (d, *J* = 8.8 Hz, 1H), 7.72 (d, *J* = 8.8 Hz, 1H), 7.59-7.50 (m, 2H), 7.45-7.38 (m, 2H), 4.42-4.35 (m, 2H), 4.07 (s, 3H), 3.85-3.65 (m, 4H), 3.53-3.45 (m, 2H), 3.38-3.35 (m, 3H), 3.20-3.10 (m, 2H), 2.97 (s, 3H), 2.46-2.37 (m, 4H), 2.15-2.00 (m, 6H). ESI-MS *m/z* 515.3 [M+H]⁺ calc. for C₃₁H₃₈N₄O₃.

***Tert*-butyl 4-[[6-methoxy-2-(5-methyl-2-furyl)-7-(3-pyrrolidin-1-ylpropoxy)-4-quinolyl]amino]piperidine-1-carboxylate (**37a**)**

To a solution of compound **22r** (2.2 g, 5.50 mmol) in 1,4-dioxane (150 mL) were added Cs₂CO₃ (3.58 g, 10.99 mmol), BINAP (342 mg, 0.55 mmol), Pd₂(dba)₃ (504 mg, 0.55 mmol) and *tert*-butyl 4-aminopiperidine-1-carboxylate (1.1 g, 5.50 mmol) in one portion at 25 °C under N₂ and the mixture was stirred at 130 °C for 5 hours under N₂. Then, the mixture was cooled to room temperature and poured into water (50 mL). The aqueous phase was extracted with EtOAc. The combined organic phase was washed with saturated brine, dried with anhydrous Na₂SO₄, filtered, concentrated under vacuum and purified by silica gel column chromatography to obtain pure compound **37a** (1.8 g, 58%) as a yellow solid. ESI-MS *m/z* 565.3 [M+H]⁺ calc. for C₃₂H₄₄N₄O₅. This intermediate was used in the next step without further characterization.

***Tert*-butyl 4-[[2-(5-ethyl-2-furyl)-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)-4-quinolyl]amino]piperidine-1-carboxylate (37b)**

To a solution of compound **33a** (622 mg, 1.50 mmol) in 1,4-dioxane (15 mL) were successively added Pd₂(dba)₃ (137 mg, 0.15 mmol), BINAP (187 mg, 0.3 mmol), Cs₂CO₃ (977 mg, 3 mmol) and *tert*-butyl 4-aminopiperidine-1-carboxylate (451 mg, 2.25 mmol) and the resulting mixture was stirred at 80 °C for 36 hours. Then, the mixture was cooled to room temperature, diluted with water and then extracted with EtOAc. The combined organic phase was washed with brine, concentrated and purified by silica gel column chromatography to obtain pure compound **37b** (950 mg, 99% crude) as a yellow solid. ESI-MS *m/z* 579.1 [M+H]⁺ calc. for C₃₃H₄₆N₄O₅. This intermediate was used in the next step without further characterization.

6-Methoxy-2-(5-methyl-2-furyl)-*N*-(4-piperidyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (38a)

A mixture of compound **37a** (1.80 g, 3.19 mmol) and HCl/EtOAc (1.0 M, 80 mL) was stirred at 25 °C for 2 hours under N₂. Then, the mixture was concentrated to dryness to give the crude product which was purified by preparative HPLC (method 2 described in supporting information) to afford pure compound **38a** (1.2 g, 85%) as yellow solid; m.p. 127-128 °C. ¹H NMR (MeOD, 400 MHz): δ 7.92 (s, 1H), 7.71 (d, *J* = 3.3 Hz, 1H), 7.57 (s, 1H), 7.17 (s, 1H), 6.46 (d, *J* = 3 Hz, 1H), 4.40 (t, *J* = 5.4 Hz, 3H), 4.12-4.05 (m, 5H), 3.85-3.75 (m, 2H), 3.60 (d, *J* = 13.1 Hz, 2H), 3.52 (t, *J* = 7.2 Hz, 2H), 3.21-3.13 (m, 2H), 2.54 (s, 3H), 2.45-2.28 (m, 4H), 2.25-2.02 (m, 6H). ESI-MS *m/z* 465.3 [M+H]⁺ calc. for C₂₇H₃₆N₄O₃.

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3 **2-(5-Ethyl-2-furyl)-6-methoxy-N-(4-piperidyl)-7-(3-pyrrolidin-1-**
4 **ylpropoxy)quinolin-4-amine (38b)**

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7 A solution of compound **37b** (868 mg, 1.50 mmol) in HCl/EtOAc (1.0 M, 40 mL) was
8 stirred at 22 °C for 3 hours. Then, the reaction mixture was concentrated to dryness to
9 give compound **38b** (850 mg, 99% crude). ESI-MS m/z 479 $[M+H]^+$ calc. for
10 $C_{28}H_{38}N_4O_3$. This intermediate was used in the next step without further
11 characterization.
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20 **N-(1-isopropyl-4-piperidyl)-6-methoxy-2-(5-methyl-2-furyl)-7-(3-pyrrolidin-1-**
21 **ylpropoxy)quinolin-4-amine (39a)**

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24 To a mixture of compound **38a** (155 mg, 0.33 mmol) and acetone (108 mg, 1.86 mmol)
25 in *i*-PrOH (40 mL) were added NaBH₃CN (117 mg, 1.86 mmol) and AcOH (112 mg,
26 1.86 mmol) in one portion at 16 °C under N₂ and the mixture was stirred at 50 °C for 15
27 hours. Then, the mixture was cooled to 16 °C, filtered and concentrated in vacuum. The
28 residue was purified by preparative HPLC (method 4 described in supporting
29 information) to obtain pure compound **39a** (61.4 mg, 37%) as a yellow solid; m.p. 106-
30 107 °C. ¹H NMR (MeOD, 400 MHz): δ 7.82 (s, 1H), 7.59 (d, J = 3.2 Hz, 1H), 7.51 (s,
31 1H), 7.09-7.07 (m, 1H), 6.43 (d, J = 2.8 Hz, 1H), 4.36-4.33 (m, 3H), 4.04 (s, 3H), 3.86-
32 3.76 (m, 2H), 3.65-3.57 (m, 3H), 3.51-3.47 (m, 2H), 3.36-3.31 (m, 2H), 3.20-3.10 (m,
33 2H), 2.50 (s, 3H), 2.45- 2.33 (m, 4H), 2.25-2.13 (m, 4H), 2.10-2.02 (m, 2H), 1.43 (d, J =
34 6.8 Hz, 6H). ESI-MS m/z 507.4 $[M+H]^+$ calc. for C₃₀H₄₂N₄O₃.
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50 **2-(5-Ethyl-2-furyl)-N-(1-isopropyl-4-piperidyl)-6-methoxy-7-(3-pyrrolidin-1-**
51 **ylpropoxy)quinolin-4-amine (39b)**
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To a mixture of compound **38b** (120 mg, 0.25 mmol) and acetone (81 mg, 1.40 mmol) in *i*-PrOH (20 mL) were added NaBH₃CN (88 mg, 1.40 mmol) and AcOH (84 mg, 1.4 mmol) in one portion at 16 °C under N₂ and the mixture was stirred at 50 °C for 15 hours. Then, the mixture was cooled to 16 °C, filtered and concentrated in vacuum. The residue was purified by preparative HPLC (method 4 described in supporting information) to obtain pure compound **39b** (32.9 mg, 25%) as a yellow solid; m.p. 105-106 °C. ¹H NMR (MeOD, 400 MHz): δ 7.83 (s, 1H), 7.60 (d, *J* = 3.6 Hz, 1H), 7.52 (s, 1H), 7.10 (s, 1H), 6.45 (d, *J* = 3.6 Hz, 1H), 4.36-4.31 (m, 3H), 4.02 (s, 3H), 3.87-3.77 (m, 2H), 3.70-3.60 (m, 3H), 3.52-3.45 (m, 2H), 3.35-3.31 (m, 2H), 3.20-3.10 (m, 2H), 2.90-2.82 (m, 2H), 2.46-2.35 (m, 4H), 2.25-2.15 (m, 4H), 2.11-2.00 (m, 2H), 1.43 (d, *J* = 6.8 Hz, 6H), 1.36 (t, *J* = 7.6 Hz, 3H). ESI-MS *m/z* 521.4 [M+H]⁺ calc. for C₃₁H₄₄N₄O₃.

***N*-(1-cyclopropyl-4-piperidyl)-6-methoxy-2-(5-methyl-2-furyl)-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (39c)**

To a solution of compound **38a** (1 g, 2.15 mmol) and (1-ethoxycyclopropoxy)-trimethyl-silane (2.09 g, 12 mmol) in MeOH (100 mL) were added NaBH₃CN (750 mg, 11.98 mmol) and AcOH (719 mg, 11.98 mmol) in one portion at 25 °C under N₂ and the mixture was stirred at 25 °C for 10 minutes. Then, the mixture was heated to 60 °C overnight. Then, the reaction mixture was concentrated and purified by preparative HPLC (method 2 described in supporting information) to obtain pure compound **39c** (333 mg, 31%) as a yellow solid; m.p. 120-122 °C. ¹H NMR (MeOD, 400 MHz): δ 7.82 (s, 1H), 7.58 (d, *J* = 3.5 Hz, 1H), 7.50 (s, 1H), 7.10 (s, 1H), 6.45 (dd, *J* = 3.5, 0.88 Hz, 1H), 4.35 (t, *J* = 5.51 Hz, 3H), 4.04 (s, 3H), 3.85-3.75 (m, 4H), 3.50 (t, *J* = 7.06 Hz, 3H), 3.21-3.11 (m, 3H), 2.85 (br s, 1H), 2.52 (s, 3H), 2.43-2.36 (m, 4H), 2.23-2.06 (m,

6H), 1.10-1.06 (m, 2H), 1.05-1.00 (m, 2H). ESI-MS m/z 505.4 $[M+H]^+$ calc. for $C_{30}H_{40}N_4O_3$.

***N*-(1-cyclopropyl-4-piperidyl)-2-(5-ethyl-2-furyl)-6-methoxy-7-(3-pyrrolidin-1-ylpropoxy)quinolin-4-amine (39d)**

To a mixture of compound **38b** (185 mg, 0.39 mmol) and (1-ethoxycyclopropoxy)-trimethyl-silane (406 mg, 2.30 mmol) in MeOH (20 mL) were added $NaBH_3CN$ (146 mg, 2.33 mmol) and AcOH (140 mg, 2.6 mmol) in one portion under N_2 and the mixture was heated to 60 °C overnight. Then, the mixture was cooled and filtered. The filtrate was concentrated in vacuum and the residue was purified by preparative HPLC (method 1 described in supporting information) to obtain pure compound **39d** (46 mg, 23%) as a yellow solid; m.p. 113-115 °C. 1H NMR (MeOD, 400 MHz): δ 7.82 (s, 1H), 7.62 (d, $J = 3.6$ Hz, 1H), 7.52 (s, 1H), 7.09 (s, 1H), 6.45 (d, $J = 3.2$ Hz, 1H), 4.83-4.79 (m, 1H), 4.39-4.31 (m, 3H), 4.04 (s, 3H), 3.87-3.78 (m, 4H), 3.55-3.45 (m, 4H), 3.19-3.10 (m, 2H), 2.90-2.80 (m, 2H), 2.45-2.35 (m, 4H), 2.27-2.15 (m, 4H), 2.12-2.02 (m, 2H), 1.36 (t, $J = 7.6$ Hz, 3H), 1.11-1.04 (m, 2H), 1.02-0.95 (m, 2H). ESI-MS m/z 519.4 $[M+H]^+$ calc. for $C_{31}H_{42}N_4O_3$.

G9a and DNMT1 docking

Compound **11** was superposed to the conformation of UNC0638 in the cocrystal structure of the G9-UNC0638-SAH complex (Protein Data Bank, PDB, entry 3RJW) with the MOE program (Chemical Computing Group, <http://www.chemcomp.com>). Then, the overlaid conformation of the compound was translated into the G9a-UNC0638-SAH crystal in order to analyze the key interactions between the ligand and the methyltransferase.

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3 The GoldSuite 5.2 program (Cambridge Crystallographic Data Centre,
4 <https://www.ccdc.cam.ac.uk/pages/Home.aspx>) was used to carry out docking of
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6 compounds to DNMT1. The crystal structure of Mouse DNMT1 bound to
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8 hemimethylated CpG DNA (PDB entry 4DA4) was chosen. In order to explore both,
9
10 different binding pockets and different binding modes, a range of docking set ups where
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12 considered with emphasis on keeping adequate volume occupancy of the different
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14 binding pockets and considering protein-ligand interactions, especially those involving
15
16 conserved catalytic residues. The docking configuration was, when adequate, validated
17
18 by reproducing the crystallographic binding mode of SAH. In the final selected set-up,
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20 the docking region used was a 20-Å sphere around the carboxylate oxygen of Glu1269.
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22 The PLP scoring function was used to rank docking poses, and protein hydrogen bond
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24 constraints for binding to carboxylate of Glu-1269 were imposed on the ligand. The top
25
26 twenty best docked structures out of 100 independent genetic algorithm runs were
27
28 retrieved and visually inspected. The high-scoring pose was finally chosen as it has a
29
30 plausible binding mode with key interactions with DNMT1 and a high degree of
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32 convergence ($\text{rmsd} < 2 \text{ \AA}$) was observed among the top three ranked poses.
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39 **Comparative analysis of the electrostatic potential**

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41 Electrostatic maps for each compound where calculated with VIDA (OpenEye,
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43 <http://www.eyesopen.com/vida>). Previously, all compounds where overlaid to the
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45 reference compound **11**, translated into the protein cavity and relaxed to adapt their
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47 conformation to this cavity using MOE.
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53 **G9a enzyme activity assay**

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3 The biochemical assay to measure G9a enzyme activity relies on time-resolved
4 fluorescence energy transfer (TR-FRET) between europium cryptate (donor) and
5 XL665 (acceptor). TR-FRET is observed when biotinylated histone monomethyl-H3K9
6 peptide is incubated with cryptate-labeled anti- H3K9me2 antibody (CisBio Cat#
7 61KB2KAE) and streptavidin XL665 (CisBio Cat#610SAXLA), after enzymatic
8 reaction of G9a.
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11 The assay was carried out during 1 hour at room temperature, in a final volume of 20
12 μ L, with 0.2 nM human G9a enzyme, 40nM biotinylated histone monomethyl-H3K9
13 peptide, 20 μ M S-adenosylmethionine (SAM) and different final concentrations of
14 tested compounds in assay buffer (50 mM Tris-HCl, 10 mM NaCl, 4 mM DTT, 0.01%
15 Tween-20 pH 9). The final percentage of DMSO was 0.5%. After incubation time
16 enzyme activity was stopped by adding 150 nM of cryptate-labeled anti-H3K9me2
17 antibody and 16 μ M of streptavidin XL665 beads. After one hour of incubation at room
18 temperature, fluorescence was measured at 620 nm and 665 nm. A ratio (665 nm / 620
19 nm) was then calculated in order to minimize medium interferences. Positive control
20 was obtained in the presence of the vehicle of the compounds. Negative control was
21 obtained in the absence of G9a enzyme activity. Calculated IC₅₀ values were determined
22 using GraphPrism using 4-parameters inhibition curve. Compounds were tested in
23 duplicate at different days, within an experimental error of 0.3 log units. If absolute
24 pIC₅₀ difference was higher than 0.3 log units, additional replicates were performed
25 until satisfying the experimental error (by discarding individual results with values
26 outside 2 MADs of the mean value).
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51 52 53 **DNMT1 enzyme activity assay** 54 55 56 57 58 59 60

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3 The biochemical assay to measure DNMT1 enzyme activity relies on time-resolved
4 fluorescence energy transfer (TR-FRET) between lumi4-Tb (donor) and d2 (acceptor)
5 using the EPIgeneous methyltransferase assay (CisBio Cat#62SAHPEB). TR-FRET is
6 observed when antibody specific to S-adenosylhomocysteine labeled with Lumi4-Tb is
7 incubated with d2-labeled S-adenosylhomocysteine. TR-FRET signal is inversely
8 proportional to the concentration of SAH, product of DNMT1 enzyme activity, in the
9 sample.
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12 The assay was carried out during 15 minutes at 37 °C, in a final volume of 20 μ L, with
13 20 nM human DNMT1 enzyme, 1 μ g/mL poly-deoxy inosine poly-deoxy cytosine (pdI-
14 pdC) DNA, 1 μ M S-adenosylmethionine (SAM) and different final concentrations of
15 tested compounds in assay buffer (50 mM Tris-HCl, 1 mM EDTA, 1 mM DTT, 0.1%
16 Triton X-100, 5% glycerol pH 7.5). The final percentage of DMSO was 0.5%. After
17 incubation time enzyme activity was stopped by adding 2 μ L of buffer one of the
18 EPIgeneous methyltransferase kit assay. After 10 minutes at room temperature, it was
19 added 4 μ L of antibody specific to S-adenosylhomocysteine labeled with Lumi4-Tb 50
20 x and 4 μ L of d2-labeled S-adenosylhomocysteine 31x both diluted in buffer two of the
21 EPIgeneous methyltransferase kit. Fluorescence was measured at 620 nm and 665 nm
22 one hour later. A ratio (665 nm / 620 nm) was then calculated in order to minimize
23 medium interferences. Positive control was obtained in the presence of the vehicle of
24 the compounds. Negative control was obtained in the absence of DNMT1 enzyme
25 activity. Calculated IC₅₀ values were determined using GraphPrism using 4-parameters
26 inhibition curve. Compounds were tested in duplicate at different days, within an
27 experimental error of 0.3 log units. If absolute pIC₅₀ difference was higher than 0.3 log
28 units, additional replicates were performed until satisfying the experimental error (by
29 discarding individual results with values outside 2 MADs of the mean value).
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Methyltransferase Selectivity Panel

Selectivity of **11** and **39c** against 14 methyltransferases (GLP, MLL1, SET7/9, SUV39H1, SUV39H2, PRMT1, PRMT3, PRMT4, PRMT5, PRMT6, PRMT8, EZH1, EZH2, SETD2) and DNMTs (DNMT1, DNMT3A, DNMT3B) was performed by BPS Bioscience (<http://www.bpsbioscience.com/index.ph>). Binding experiments were performed in duplicate at each concentration. GLP IC₅₀ determination for **11** and **39c** was carried out at Eurofins (<https://www.eurofins.com/>) in duplicate.

Cell culture

CEMO-1, LAL-CUN-2 and MV4-11 cell lines were cultured with RPMI 1640 medium supplemented with 20% fetal bovine serum and OCI-Ly10 and OCI-Ly3 cells with IMDM supplemented with 20% human serum and 55 μ M of β -mercaptoethanol. All cell lines were maintained in culture at 37 °C in a humid atmosphere containing 5% CO₂ and were tested for mycoplasma.

Cell Proliferation Assay – MTS

Cell proliferation was analyzed after 48 hours of *in vitro* treatment using the CellTiter 96 Aqueous One Solution Cell Proliferation Assay (Promega). This is a colorimetric method for determining the number of viable cells in proliferation. For the assay, cells were cultured by triplicate at a density of 1×10^6 cells/mL in 96-well plates (100.000 cells/well, 100 μ L/well), except for OCI-Ly3 and OCI-Ly10 cell lines which were cultured at a density of 0.5×10^6 cells/mL (50.000 cells/well, 100 μ L/well). In all cases, only the 60 inner wells were used to avoid any border effects. After 48 hours of

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3 treatment, plates were centrifuged at 800g for 10 minutes and medium was removed.
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5 Then, cells were incubated with 100 μL /well of medium and 20 μL /well of CellTiter 96
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7 Aqueous One Solution reagent. After 1-3 hours of incubation at 37 $^{\circ}\text{C}$, absorbance was
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9 measured at 490 nm in a 96-well plate reader. The background absorbance was
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11 measured in wells with only cell line medium and solution reagent. Data was calculated
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13 as a percentage of total absorbance of treated cell / absorbance of non-treated cells.
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19 **Cytotoxicity in THLE-2 cells**

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23 Cytotoxic effects of assayed compounds were tested using the immortalized human
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25 liver cell line THLE-2 (ATCC CRL-2706), cultured in BEGM medium (Clonetics #CC-
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27 4175). Medium was completed by adding 0.7 $\mu\text{g}/\text{mL}$ phosphoethanolamine, 0.5 ng/mL
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29 epidermal growth factor, antibiotics (penicillin and streptomycin) and 10% fetal bovine
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31 serum (FBS). Cells were plated in 96-well black microplates at 10,000 cells/well and
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33 incubated at 37 $^{\circ}\text{C}$ (5% CO_2 , 95% humidity) for 24 h. Test compounds were solubilized
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35 in 100% DMSO and then diluted with cell culture medium containing 10% DMSO. The
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37 final concentrations of the test compounds (1% DMSO) ranged from 0-100 μM in a final
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39 volume of 200 μL . After 72 h, cell viability in each well was determined by measuring
40
41 the concentration of cellular adenosine triphosphate (ATP) using the VialightTM Plus
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43 Cell Proliferation/Cytotoxicity Kit as described by the manufacturer (Cambrex, East
44
45 Rutherford, NJ). After addition of cell lysis buffer, test plate was incubated for 45 min
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47 at room temperature (orbital shaker). ATP monitoring solution was added and ATP
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49 concentration determined by reading luminescence using a Envision plate reader
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51 (PerkinElmer). The percentage of viable cells relative to the non-drug treated controls
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53 was determined for each well and LC_{50} values were calculated as concentrations
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3 projected to kill 50% of the cells following a 72 h exposure. Results are the average of
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5 at least two independent experiments.
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10 **Western blot**

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12 After 48 hours of treatment, CEMO-1 cells were washed twice with PBS, being the last
13 centrifugation of 4000 rpm for 10 min at 4 °C. Histone extraction was performed as
14 recommended by Upstate Biotechnology. Briefly, cells were homogenized in 5 volumes
15 of lysis buffer (10 mM HEPES, pH 7.9; 1.5 mM MgCl₂; 10 mM KCl; 0.5 mM DTT;
16 protease inhibitor cocktail (Complete Mini, Cat No 11836153301, Roche) and HCl was
17 added to a final concentration of 0.2 M. After incubation on ice for 30 min, the
18 homogenate was centrifuged at 11000 g for 10 min at 4 °C, and the supernatant was first
19 dialyzed twice against 0.1 M glacial acetic acid (1 hour each time) and then three times
20 against water for 1 hour, 3 hours and overnight, respectively. The histone concentration
21 in the extract was measured using the dye-binding assay of Bradford. 10 µg of histone
22 was separated on 15 % SDS-PAGE gel and transferred to a nitrocellulose membrane.
23
24 The membrane, after being blocked with Tropix I-block blocking reagent (Cat No
25 AI300, Tropix) in PBS with 0.1 % of Tween-20 and 0.02 NaN₃, was incubated with the
26 primary antibody against H3K9me2 (Mouse monoclonal antibody to Histone H3
27 dimethyl K9, Cat No ab1220, Abcam) diluted 1:2000 overnight at 4 °C and then with
28 alkaline phosphatase-conjugated secondary antibodies. Bound antibodies were revealed
29 by a chemiluminiscent reagent (Tropix) and detected using HyperfilmTM enhanced
30 chemiluminescence. Total H3 was used as a loading control (diluted 1:50000 overnight
31 at 4 °C or for 1 hour at rt) (Anti-Histone H3, CT, pan, rabbit polyclonal, Cat No 07-690,
32 Millipore).
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Liquid chromatography tandem-mass spectrometry (LC-MS/MS)

After 48 h of treatment, cells were washed twice with PBS and genomic DNA was extracted using a DNA kit (Nucleo Spin Tissue, Cat No 74095250, Macherey-Nagel) following the manufacturer's instructions. DNA purity and concentration was measured using a NanoDrop spectrophotometer (Thermo Scientific). For the analysis of methylation by mass-spectrometry, 1 μ g of genomic DNA was digested with DNA degradase plus (ZYMO RESEARCH) and subjected to mass spectrometry (liquid chromatography electrospray ionization tandem mass spectrometry). All samples were analyzed using an Agilent 1200 liquid chromatograph (Agilent Technologies, Wilmington, DE, USA). An Agilent Zorbax Eclipse XDB-C18 column (2.1 \times 150 mm, 3.5 μ m particle size, Agilent Technologies, Wilmington, DE, USA) was used.

DNA methylation analysis with Human Methylation array

After 48 h of treatment, cells were washed twice with PBS and genomic DNA was extracted using a DNA kit (Nucleo Spin Tissue, Cat No 74095250, Macherey-Nagel) following the manufacturer's instructions. DNA purity and concentration was measured using a NanoDrop spectrophotometer (Thermo Scientific). 500ng of genomic DNA of MV4-11 control cell lines and these cells treated with 10nM and 50nM of compound **39c** were bisulfite converted. Converted DNA was hybridized onto the MethylationEPIC BeadChip (Illumina), which interrogate over 850,000 methylation sites quantitatively across the genome at single-nucleotide resolution, as previously described.⁷³

CYP Inhibition

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3 The inhibitory effect of the compounds on five human cytochrome P450s (1A2, 2C9,
4 2C19, 2D6, and 3A4) was evaluated in human liver microsomes at WuXi
5 (<http://www.wuxi.com/>). Compounds were prepared at 10 μM , and the corresponding
6 substrates for each P450 isoform (20 μL) were incubated with 140 μL of liver
7 microsomes (0.286 mg/mL; BD Gentest) and NADPH cofactor (20 μL , 1 mM) for 10
8 min at 37 $^{\circ}\text{C}$. The reaction was terminated by adding 400 μL of cold stop solution (200
9 ng/mL tolbutamide in ACN), and samples were centrifuged at 1500g for 20min.
10 Supernatants were analyzed by LC-MS/MS (Shimadzu LC 20-AD-API 4000) using the
11 peak area ratio of the analyte/internal standard. Compounds and positive controls were
12 tested in duplicate. The percentage of inhibition was calculated as the ratio of substrate
13 metabolite detected in treated and non-treated wells.
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29 **Metabolic Stability**

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31 Test compounds (1 μM , 5% MeOH in potassium phosphate buffer) were incubated with
32 human (catalogue no. 452161 from BD Gentest) and mouse (catalogue no. M1000,
33 Xenotech) liver microsomes at 37 $^{\circ}\text{C}$ for 10 min. Liver microsomes were at a final
34 assay concentration of 0.7 mg protein/mL. The reaction was started by the addition of
35 90 μL of NADP cofactor solution and stopped by the addition of 300 μL of stop
36 solution (ACN at 4 $^{\circ}\text{C}$, including 100 ng/mL tolbutamide as an internal standard) after
37 20 min of incubation. The samples were shaken for 5 min and then centrifuged for 20
38 min at 1500g. A 100 μL aliquot of the supernatant was transferred to eight new 96-well
39 plates with 300 μL of HPLC water and centrifuged at 1500g for LC-MS/MS analysis
40 (Shimadzu LC 10-AD-API 4000). An injection volume of 10 μL was added to a
41 Phenomenex Synergi C18 column eluting with formic acid in water or ACN at a flow
42 rate of 800 $\mu\text{L}/\text{min}$. The percent loss of parent compound was calculated from the peak
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area ratio of the analyte/internal standard. Compounds and positive controls were tested in duplicate.

PAMPA Permeability

The permeability of compounds was evaluated with the parallel artificial membrane permeation assay (PAMPA) as an *in vitro* model of passive diffusion. Donor solutions of test compounds (180 μ L, 50 μ M in PBS/EtOH 70:30) were added to each well of the donor plate, whose PVDF membrane was precoated with 4 μ L of a 20 $\text{mg}\times\text{mL}^{-1}$ PBL/dodecane mixture. PBS/EtOH (180 μ L) was added to each well of the PTFE acceptor plate. The donor and acceptor plates were combined together and incubated for 18 h at 20 $^{\circ}$ C without shaking. In each plate, compounds and controls were tested in duplicate. Drug concentration in the acceptor, the donor, and the reference wells was determined using the UV plate reader with 130 μ L of acceptor and donor samples. Permeability rates (P_e in nm s^{-1}) were calculated with Equation (1). The permeability rate of each compound is the averaged value of three independent measurements.

$$\text{Equation (1) } P_e = C \times \left(-\ln \left(1 - \frac{[\text{drug}]_{\text{acceptor}}}{[\text{drug}]_{\text{equilibrium}}} \right) \right) \times 10^7 ;$$

$$\text{here } C = \frac{V_D \times V_A}{(V_D + V_A) \times \text{Area} \times \text{time}} ; V_D = 0.18 \text{ mL}; V_A = 0.18 \text{ mL}; \text{Area} = 0.32 \text{ cm}^2; \text{time} = 64800 \text{ s}; D_F = 180/130; [\text{drug}]_{\text{equilibrium}} = ([\text{drug}]_{\text{donor}} \times V_D + [\text{drug}]_{\text{acceptor}} \times V_A) / (V_D + V_A); [\text{drug}]_{\text{donor}} = (A_a/A_i * D_F)_{\text{donor}}; [\text{drug}]_{\text{acceptor}} = (A_a/A_i * D_F)_{\text{acceptor}}; A_a \text{ donor} = \text{Abs}_{\text{donor}} - \text{Abs}_{\text{vehicle}}; A_a \text{ acceptor} = \text{Abs}_{\text{acceptor}} - \text{Abs}_{\text{vehicle}}, A_i = \text{Abs}_{\text{withoutPBL}} - \text{Abs}_{\text{vehicle}}.$$

Quantification of compounds in Plasma Samples (PK studies).

In vivo PK studies had previous approval from the Animal Care and Ethics Committee of the University of Navarra (protocol numbers: 158-12 and 009-16). Compounds were

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3 measured in plasma samples using an Acquity UPLC (Waters, Manchester, UK)
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5 coupled to Xevo-TQ MS triple quadrupole mass spectrometer with an electrospray
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7 ionization (ESI) source. Compound solutions were prepared by dissolving the solid in
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9 saline and drug dosages were administered as a single intravenous injection. Blood was
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11 collected at predetermined times into EDTA-containing tubes and plasma was obtained
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13 via centrifugation (4 °C, 2500 rpm, 5 min) and stored at -80° C until analysis.
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18 Chromatographic separation was performed by gradient elution at 0.6 mL/min using an
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20 Acquity UPLC BEH C18 column (50 x 2.1 mm, 1.7 µm particle size; Waters). Mobile
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22 phase consisted of A, water with 0.1% formic acid, and B, methanol with 0.08% formic
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24 acid. Autosampler temperature was set at 7 °C and column temperature at 40 °C. For
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26 detection and quantification, the electrospray ionization operated in the positive mode,
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28 and the collision gas used was ultra-pure argon at a flow rate of 0.15 mL min⁻¹. The
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30 compound was detected using multiple reaction monitoring (MRM).
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33 Quantification was achieved by external calibration using matrix-matched standards.
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35 Concentrations were calculated using a weighted least-squares linear regression ($W =$
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37 $1/x$). Calibration standards were prepared by adding the appropriate volume of diluted
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39 solutions of the compound (made in a mixture of methanol and water, 50:50, v:v) to
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41 aliquots of 25 µL of blank plasma. 3% formic acid in methanol was added to precipitate
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43 the proteins. The mixture was then agitated for 10 min and centrifuged at 13200 rpm for
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45 20 min at 4° C. A 5 µL aliquot of the resulting supernatant was injected onto the LC-
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47 MS/MS system for analysis. Frozen plasma samples were thawed at room temperature,
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49 vortexed thoroughly, and subjected to the above described extraction procedure. PK
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51 parameters were obtained by fitting the blood concentration-time data to a non-
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53 compartmental model with the WinNonlin software (Pharsight, Mountain View, CA).
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Parameters estimated included area under the curve (AUC), half-life of the product ($t_{1/2}$), clearance (Cl), and volume of distribution (V_{ss}).

***In vivo* experiments**

All animals studies had previous approval from the Animal Care and Ethics Committee of the University of Navarra (protocol number: 041-15). For the human subcutaneous MV4-11 AML model, 10×10^6 MV4-11 cells diluted in 100 μ L of saline solution were subcutaneously inoculated in the back left flank of BALB/cA- Rag2^{-/-} γ c^{-/-} mice between 6 and 8 weeks of age (n=12). The treatment with 2.5 mg/kg (i.v.) of compound **12** was started 1 day after cell inoculation and was administered daily during 21 consecutive days (n=6). The control group (n=6) received only saline solution (diluent of compound **12**). Tumor size was analyzed every 5 days using the following method: $V \approx \frac{1}{4} D \cdot d^2$, where D and d corresponding to the longest and shorter diameter, respectively. Mice were killed 23 days after cell inoculation.

Interference compound assessment

Potential PAINS (Pan Assay Interference Compounds) liability of reported compounds was assessed according to the structural filters defined by Baell & Holloway⁷⁴ (Charts 3 and 4 and Tables 1-3 of this reference) using a customized Pipeline Pilot protocol.⁶⁵ No compound reported in the manuscript matches any of these substructures.

ASSOCIATED CONTENT

SUPPORTING INFORMATION

Protocols for preparative HPLC purification methods (S1)

Synthesis of compound **40** (S2)

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3 Synthesis of intermediates 1 to 3 (S3)

4
5 Method for High Resolution Mass Spectrometry of final compounds (S4)

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7 Methods for LCMS, analytical HPLC and UHPLC (S5)

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9 HRMS and purities of final compounds (S6)

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11 HPLC traces of final compounds (S7)

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13 Superposition of the autoinhibited crystal structure of hDNMT1 against the productive
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15 crystal structure of mDNMT1 (S8)

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17 Binding mode of compound **23m** into G9a (S9)

18
19 Competition experiments performed with compound **23m** (S10)

20
21 Results of MethylationEPIC BeadChip assay (S11)

22
23 Selectivity profiling of compounds **11** and **39c** against a panel of 14 lysine and arginine
24
25 methyltransferases and DNMTs (S12)

26
27 Plasmatic concentrations of compounds **34a** (S13) and **39c** (S14)

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29 Molecular formula strings and some data.

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31 This material is available free of charge via the Internet at <http://pubs.acs.org>.

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35 **PDB ID Codes:**

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37 **2**, 3K5K; **3**, 3RJW; 4DA4

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51 **Notes**

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53 These authors declare no competing financial interest.

ACKNOWLEDGEMENTS

We thank the Foundation for Applied Medical Research (FIMA), University of Navarra (Pamplona, Spain), Asociación de Amigos of University of Navarra as well as to Fundación Fuentes Dutor for financial support. This work has been partially supported through Ministerio de Economía y Competitividad and grants from Instituto de Salud Carlos III PI10/01691, PI13/00862, PI13/01469, PI14/01867, PI16/02024, PI17/00701, RTICC RD12/0036/0068, CIBERONC (CB16/12/00489) (Co-finance with FEDER funds), ERA-NET programs TRANSCAN-2 JTC EPICA by the “Torres Quevedo” Subprogram (PTQ-11-04777 and PTQ-14-07320 I.D.M), Fundacio´ La Marato´ de TV3 (20132130-31-32), Gobierno de Navarra (40/2016, PT053/2016 and PT027/2017) and FSE (Inncorpora-Torres Quevedo grant), Plan Nacional de I+D+I 2013-2016/FEDER (PI15/00892 to M.F.F.), the Asturias Regional Government (GRUPIN14-052 to M.F.F.) and the IUOPA-Obra Social Liberbank-Cajastur. Finally, we would like to thank OpenEye Scientific Software for providing us with an academic license for use of its software.

ABBREVIATIONS

ACN; acetonitrile; AcOH, acetic acid; ADME, absorption, distribution, metabolism and excretion; ALL; acute lymphoblastic leukemia; AML; acute myeloid leukemia; AUC, area under the curve; BINAP, 2,2'-bis(diphenylphosphino)-1,1'-binaphthyl; BOC, *tert*-butoxycarbonyl; BPO, benzoyl peroxide; CDI, 1,1-carbonildiimidazol; Cl, clearance; Cpd, compound; DEAD, diisopropyl azodicarboxylate; DIEA, N,N-Diisopropylethylamine; DLBCL, diffuse large B-cell lymphoma; DME, dimethoxyethane; DMF, dimethylformamide; DMSO, dimethylsulfoxide; DNA, deoxyribonucleic Acid; DNMT, DNA methyltransferase; DTT, *dithiothreitol*; EDTA,

1
2 ethylenediaminetetraacetic acid; EHMT2, euchromatic histone methyltransferase 2;
3
4 ESI-MS, electrospray ionisation mass spectrometry, Et₃N, triethylamine; EtOAc, ethyl
5
6 acetate; EtOH, ethanol; FDA, food and drug administration; HDAC, histone
7
8 deacetylase; HEPES, 4-(2-Hydroxyethyl)piperazine-1-ethanesulfonic acid; HIV, human
9
10 immunodeficiency virus; HLM, human liver microsomes; HPLC, High-performance
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12 liquid chromatography; *i*-PrOH, propan-2-ol; iPSCs; inducible pluripotent stem cells;
13
14 i.v., intravenous; KMT1C, lysine methyltransferase 1C; LCMS, liquid chromatography–
15
16 mass spectrometry, MDR, multidrug resistance; MeOH, methanol; MLM, mouse liver
17
18 microsome; m.p., melting point; MRM, multiple reaction monitoring; MST, microscale
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20 thermophoresis; MW, microwave; NADPH, nicotinamide adenine dinucleotide
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22 phosphate hydrogen; NaOMe, sodium methoxide; NMR, nuclear magnetic resonance;
23
24 PAMPA, parallel artificial membrane permeability assay; PBS, *phosphate buffered*
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26 saline; Pd₂(dba)₃, tris(dibenzylideneacetone)dipalladium(0); PDB, protein data bank;
27
28 pdI-pdC, poly-deoxy inosine poly-deoxy cytosine; Ph, phenyl; PK, pharmacokinetic;
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30 PMT, protein methyltransferase; PTFE, polytetrafluoroethylene; PVDF, polyvinylidene
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32 difluoride; RNA; ribonucleic acid; rt, room temperature; Rt, retention time; SAH, S-
33
34 adenosyl-L-homocysteine; SAM, S-adenosyl methionine; SAR, structure-activity
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36 relationship; *t*_{1/2}, half-life of the product; *t*-BuOK, potassium *tert*-butoxide; THF,
37
38 tetrahydrofuran; TLC, thin layer chromatography; TMS, tetramethylsilane; TR-FRET,
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40 time-resolved fluorescence resonance energy transfer; UPLC, ultra performance liquid
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42 chromatography; UV, ultraviolet; V_{ss}, volume of distribution; xantphos, 4,5-
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44 bis(diphenylphosphino)-9,9-dimethylxanthene.
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Table of Contents Graphic

